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| <b>(54) Title:</b> A METHOD OF INDUCING A CTL RESPONSE   |  |   |  |
| <b>(57) Abstract</b><br><br>A method of inducing a cytotoxic T-lymphocyte (CTL) response to an antigen is disclosed. The method involves delivering the antigen to the lymphatic system of an animal regularly over a sustained period of time using, e.g., an osmotic pump. The method is advantageous over prior art methods for inducing a CTL response in that it does not require repetitive immunizations or the use of adjuvants. The method of the present invention can be used for the induction of CTLs in tumor or infectious disease immunotherapy. |  |   |  |

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## A METHOD OF INDUCING A CTL RESPONSE

### CROSS REFERENCE

5 This application is a continuation-in-part of U.S. Patent Application 08/988,320 filed December 10, 1997.

### FIELD OF THE INVENTION

10 The invention relates to a method of inducing a CTL response to an antigen by sustained, regular delivery of the antigen to an animal so that the antigen reaches the lymphatic system.

### BACKGROUND OF THE INVENTION

15 Cytotoxic T lymphocytes (CTL) are white blood cells found in the blood, spleen and lymph. CTL have the ability to attack and kill other cells of the body in a highly specific manner. When CTL are stimulated by specific antigen, they migrate through the tissues of the body on a "search and destroy" mission for cells bearing the specific antigen. Whether of viral origin or tumor associated, CTL detect antigen that is bound to major histocompatibility complexes (MHC) on the surface of potential target cells. Once CTL have identified the antigen on the cell surface, their function is to deliver a lethal hit to the cell.

20 Although there are hundreds of millions of CTL that reside in the spleen, each individual CTL exclusively responds to a unique and specific antigen. These individual CTL, dubbed CTL precursors (CTLp), undergo cell division or proliferate upon activation by specific antigen to produce daughter cells with precisely the same antigen specificity as the parent cell. This proliferation increases the total number, and thus the frequency, of that specific CTLp in the body. A proportion of these newly generated CTL briefly recirculate through the body (termed effector CTL), and have the ability to identify and destroy cells bearing the specific antigen which they recognize. A significant body of experimental evidence suggests that CTL specific for tumor antigens can inhibit tumor growth. Unfortunately, most tumors have only a very weak capacity to stimulate CTL responses and there has been no means of inducing a CTL response then sustaining it over a period of time sufficient to continuously inhibit tumor growth. While many attempts to directly increase the capacity of tumor cells to stimulate tumor-clearing CTL responses in patients have been made, such attempts have met with limited success. Technical advances over

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the past ten years have, however, enabled the identification of natural peptide antigens that are present on tumor cells and which are recognized by CTL. These antigen targets include proteins expressed in significant overabundance, abnormally expressed embryonic proteins, protein products from mutated oncogenes or suppressor genes, or proteins derived from cancer-causing viruses present in tumor cells. The challenge has been to find a way in which to administer an antigen so that it induces an anti-tumor CTL response and maintains it over time. While many attempts have now been made to use these antigens clinically in a vaccine, the results have been less than satisfactory.

An explanation of why CTL therapies have been largely ineffective at eradicating or controlling tumors in a clinical setting include the following:

- (a) Vaccine designs have been inadequate at initiating strong CTL responses;
- (b) Tumor cells can down regulate MHC molecules, resulting in the loss of antigen presentation from the surface of cells, thereby escaping detection by CTL;

- (c) After induction, effector CTL recirculation through the body is highly transient;

- (d) After recirculation, CTL return to the spleen where they reside in a non-active or resting state, and an increase in the numbers of CTLp residing in the spleen does not reflect active CTL immunity;

- (e) In the case of tumors, regrowth of residual tumor cells following immunization goes undetected by CTLp residing in spleen in a "resting" state;

- (f) Because CTL-stimulating antigen-presenting cells (APC) are targeted for destruction by the same CTL that they have activated, the CTL response is self-limiting, which precludes, under normal circumstances, the continuous stimulation for a long-lived CTL response.

A growing repertoire of tumor associated antigens are being discovered that are recognized by CTL. A variety of techniques have been suggested to render these antigens effective in CTL vaccines. These include immunization using synthetic peptide antigens mixed with an immunostimulatory adjuvant, such as the bacterial toxin BCG; immunization with multiple antigenic peptide systems (MAPS); immunization with "professional" antigen presenting cells, which are isolated from the patient, pulsed with peptide antigen and inoculated back into the patient as a vaccine; immunization with peptides designed to stimulate both CTL and T helper cell populations; immunization with viruses or bacteria engineered to express tumor



antigens; and immunization with polynucleotide expression vectors (so called DNA vaccines). Unfortunately, none of these approaches has been an unqualified success. As discussed above, the lack of vigorous therapeutic effects with these vaccine platforms reflects at least to some degree problems associated with inducing a strong initial CTL response and with maintaining ongoing "active" CTL immunity.

Studies by Glenny during the first quarter of the century revealed that aluminum compounds could enhance the strength of diphtheria vaccines. This was ostensibly the first of a long history of observations supporting a "depot" theory of immunization, which postulates that antigen slowly leaking into the tissues over an extended time correlates with the antigenic potency of a vaccine. Today, this antigen depot paradigm forms the intellectual backdrop to most adjuvant development programs. In one form or another, depot type adjuvants are intended to prolong the course of antigen delivery, by forming a lesion at the site of injection, or simply by the slow degradability of the adjuvant itself, which mixed with the specific antigen forms a depot at the site of injection. A second function generally attributed to adjuvants are their immunostimulatory effects, which appears to trigger the immune system to respond to the vaccine. However, adjuvants are a double-edged sword. They have inherent toxicities. But it is a feature of these toxicities that achieves a desired immunostimulatory and/or depot effect. Side effects such as tissue damage and granulomatous reaction at the site of injection, fever, and in some cases systemic reactions, such as Reiter's syndrome-like symptoms, uveitis and arthritis, are some of the risks associated with the use of adjuvants. Currently, the only adjuvant approved by the FDA is alum. It is relatively safe but does have side effects such as erythema, subcutaneous nodules, contact hypersensitivity, and granulomatous inflammation. More importantly, alum only acts to potentiate a limited number of antigens, and it very predominantly stimulates humoral antibody responses rather than CTL immunity. Thus so far adjuvants have proved to be very ineffective components for vaccines aimed at inducing clinically relevant CTL responses.

Recent attempts to induce CTL responses using dendritic cells or other antigen presenting cells, despite being cumbersome, have shown some promise. New recombinant virus or bacterial systems carrying genes for specific antigen are effective at inducing primary CTL responses. The most effective viruses, for example, that induce strong CTL responses are those which replicate aggressively in the host. Yet because of the risk for serious or lethal complications as a result of

infection, recombinant virus used in a cancer vaccine must be only weakly replicative, or be completely replication deficient. This trade-off between virulence and efficacy is at present an intractable problem.

5 DNA (or polynucleotide) vaccines are also being developed for the purpose of inducing CTL immunity. Once again, the system has intrinsic limitations that preclude its efficacy in inducing long-lasting CTL immunity. The DNA vaccines consist of a plasmid or similar genetic construct for expressing the antigen of interest. Uptake of the plasmid system by cells of the body results in expression of the antigen and induction of CTL. However, once cells expressing the construct have succeeded  
10 in inducing CTL, they are themselves targets for eradication by the CTL. The CTL inducing effect is thus again transient. Moreover, the polynucleotide vaccines have thus far suffered from poor efficiency in terms of CTL induction.

With difficulties in achieving strong primary and/or persisting CTL responses, there are a number of clinical trial groups now using repeated injections of cancer  
15 vaccines. The use of antigenically complex materials in the vaccine formulation, such as recombinant virus, or the costs associated with repetitive treatment using cultured APC will, however, make such an approach difficult. On the one hand, repetitive immunization with antigenically complex materials drives the immune system to elaborate a humoral antibody, as opposed to a CTL response, while on the other hand,  
20 use of a minimal CTL antigen (such as a nonamer peptide) which does not efficiently drive an antibody response, has also failed to induce a CTL response. Attempts to develop adjuvants that enhance the immunostimulatory aspects of minimal CTL antigens have resulted in the production of materials (i.e. adjuvants) that also induce a competing humoral immune response, or, which simply offer little CTL stimulatory  
25 effect.

It has also been suggested that certain controlled release technology using microspheres or liposomes with subunit antigens and peptides might be effective to enhance immunogenicity. The combination of sustained release and depot effect is suggested to reduce the amount of antigen needed and eliminate booster shots.  
30 However, the preparation of such compositions is difficult and unpredictable, and vaccine formulations based on this technology have not been translated into effective clinical treatments.

As can be seen from the foregoing, there has been little success at developing a CTL vaccine that is both capable of inducing a strong CTL response then sustaining

that response over time. The development of a vaccine with these capabilities is essential before effective anti-tumor therapy based on CTL immunity can be contemplated.

### **OBJECTS OF THE INVENTION**

5           An object of this invention is to provide a method for inducing or sustaining a specific CTL immunological response in a mammal over time.

          Another object of this invention is to provide a method for treating a mammal having a malignant tumor or infectious disease by inducing and sustaining an immunological attack on the malignant tumor or infectious disease in the mammal.

10           It is a further object of this invention to provide an article of manufacture useful for inducing and sustaining a specific immunological CTL response in a mammal over time.

          It is a further object of this invention to provide an article of manufacture useful for treating a mammal having a malignant tumor or infectious disease, which  
15   article is designed to induce and maintain an immunological attack on the malignant tumor or infectious disease in the mammal.

          It is a further object of this invention to provide a portable device for sustained delivery of an antigen to a mammal having a malignant tumor or infectious disease, where the antigen stimulates the mammal's immune system to attack the tumor or  
20   infectious disease and the device is located outside the mammal.

          It is still a further object of this invention to provide an implantable device for sustained delivery of an antigen to a mammal having a malignant tumor or infectious disease, where the antigen stimulates the mammal's immune system to attack the tumor or infectious disease.

25           It is a further object of this invention to provide antigen compositions and containers therefor that are useful in the methods, devices, and/or articles of manufacture of this invention.

          Other objects of this invention may be apparent to those of skill in the art by reading the following specification and claims.

### **SUMMARY OF THE INVENTION**

30           In one aspect of the invention, a method is provided for inducing an immunological CTL response to an antigen by sustained, regular delivery of the antigen to a mammal so that the antigen reaches the lymphatic system. In particular, the antigen is delivered to the mammal at a level sufficient to induce an immunologic

CTL response in the mammal and the level of the antigen in the mammal's lymphatic system is maintained over time sufficient to maintain the immunologic CTL response. Preferably, the antigen is delivered directly to the mammal's lymphatic system, such as to the spleen, a lymph node or lymph vessel.

5 Also provided is a method of treating an animal having a disease, or being predisposed to a disease, to which the animal's immune system mounts a cell-mediated response to a disease-related antigen to attack the disease. In this aspect of the invention, a disease-matched antigen is delivered to the animal at a level sufficient to induce an increased CTL-response in the animal which is then maintained in the  
10 animal by sustained, regular delivery of the disease-matched antigen to the animal for a time sufficient to treat the disease. The sustained, regular delivery of the antigen is done in a manner that maintains the level of antigen in the animal's lymphatic system. Preferably, the sustained, regular delivery is achieved by pumping a physiologically-acceptable, composition of the antigen from a device held external of or implanted in  
15 the animal's body so that the antigen reaches the animal's lymph system. Optionally, a cytokine that is capable of enhancing the CTL response is delivered and/or maintained along with the antigen. Diseases addressed in this manner include cancer and pathogenic diseases.

In a further aspect of the invention, an article of manufacture is provided for  
20 delivering an antigen that induces a CTL response in an animal. In particular, the article comprises a reservoir of a physiologically-acceptable, antigen-containing composition that is capable of inducing a CTL response in an animal; a pump connected to the reservoir to deliver the composition at a defined rate; a transmission line to discharge the composition from the reservoir; and, optionally, a delivery line  
25 connected to the transmission line, which delivery line is of a size suitable for positioning in the animal and for delivery of the composition in a manner that reaches the lymphatic system of the animal.

In a further aspect of the invention, a process is provided for preparing a system useful for inducing a sustained CTL response in an animal needing such a  
30 response, which comprises placing a physiologically-acceptable, antigen-containing composition in a reservoir having a pump for delivering the composition at a defined rate through a transmission line to the animal.

**BRIEF DESCRIPTION OF THE DRAWINGS**

The invention will now be described in relation to the drawings in which:

Figure 1 is a graph showing the lysis of target cells by CTL versus the effector/target ratio when antigen is delivered as a single dose (circles) and when antigen is delivered by a continuous pump (triangles).

Figure 2 (A and B) are graphs showing the lysis of target cells by CTL versus the effector/target ratio when antigen is delivered as a single dose (circles), when antigen is delivered by a continuous pump (triangles) and negative control (squares) at (A) 36 hours and (B) 7 days.

Figure 2C is a graph showing the footpad swelling versus time when antigen is delivered as a single dose (circles) and when antigen is delivered by a continuous pump (triangles).

Figure 3 is a graph showing the lysis of target cells by CTL versus the dose of the peptide antigen when the antigen is delivered subcutaneously, intravenously and intrasplenically.

Figure 4 is a bar graph showing tritiated thymidine uptake in CTL cells induced by antigen introduced intravenously, intrasplenically and subcutaneously.

Figure 5 is a rough schematic of a human lymphatic system.

**DETAILED DESCRIPTION OF THE INVENTION****Method of treatment**

One aspect of this invention is a method for inducing or sustaining a specific immunological response (*i.e.*, a CTL response) in an animal that has a disease (or predisposition to a disease) in which the animal's immune system may attack the disease with a natural CTL response. The response and diseases are discussed in greater detail hereinafter. The method has particular value for treating an animal having a malignant tumor in order to inhibit the growth of the tumor or for treating a chronic infectious disease such as hepatitis or AIDS.

The method, along with other aspects of the invention, is useful in an animal having an immune system that includes a lymphatic system. This generally includes vertebrates, specifically mammals and particularly humans. Thus, this invention will find use in treating humans of all ages as well as in treating animals, *i.e.* in veterinary uses. The invention may be used for treating livestock such as cattle, sheep, pigs, goats, and the like or for treating household pets such as dogs, cats, rabbits, hamsters, mice, rats, and the like. The primary use will be for treating humans that are in need

of having a specific immunological response sustained for treatment of a disease such as cancer or chronic infections.

A key aspect of this invention is the delivery of an appropriate antigen to the lymphatic system of the animal being treated and sustaining the delivery over time.

5 This is based in part on the observation that a strong induction and a sustained CTL response require ongoing antigenic stimulation of the lymphatic system. In a human, the lymphatic system includes lymph, lymphocytes, lymph vessels, lymph nodes, tonsils, the spleen, the thymus gland, and bone marrow. The lymphatic system performs three basic functions. First, it helps maintain fluid balance in the tissues.  
10 Approximately 30 L of fluid pass from the blood capillaries into the interstitial spaces each day, whereas only 27 L pass from the interstitial spaces back into the blood capillaries. If the extra 3 L of interstitial fluid were to remain in the interstitial spaces, edema would result, causing tissue damage and eventual death. These 3 L of fluid (*i.e.* lymph) enter the lymph capillaries, then passes through the lymph vessels to  
15 return to the blood. Lymph is similar in composition to plasma. In addition to water, lymph contains solutes derived from two sources: (1) substances in plasma such as ions, nutrients, gases, and some proteins pass from blood capillaries into the interstitial spaces to become part of the lymph; and (2) substances derived from cells within the tissues such as hormones, enzymes, and waste products are also found in  
20 lymph.

The lymphatic system's second basic function is to absorb fats and other substances from the digestive tract. Special lymph vessels called lacteals are in the lining of the small intestine. Fats enter into the lacteals and pass through the lymph vessels to the venous circulation. The lymph passing through these capillaries has a  
25 milky appearance because of its fat content, and it is called chyle.

The third basic function of the lymphatic system is to act as part of the body's defense system. The lymph nodes filter lymph, and the spleen filters blood, removing microorganisms and other foreign substances. This third function is the function most important to this invention in that the antigen must be delivered to the lymph system  
30 at a level sufficient to elicit the desired, specific immunological response in the animal. Figure 5 is a schematic representation of a human lymphatic system showing the major lymphatic organs and vessels.

As hereinbefore mentioned, the present invention relates to a method of inducing or sustaining a specific immunological response (particularly a CTL

response) to an antigen in an animal over time. The method comprises delivering the antigen to the animal in a manner that delivers the antigen into the lymphatic system of an animal to sustain the desired response over time. Generally this is done by establishing a mechanism to transfer an antigen from a reservoir to the animal's lymphatic system on a regular basis over time. The antigen may be delivered by a variety of methods that target intralymphatic presentation, including subcutaneous injection, direct injection into the lymphatic system by an antigen delivery vehicle that is implanted, preferably at or near a lymphatic organ, or by an antigen delivery vehicle that is external to the animal but contains a means (e.g. a needle or catheter) to deliver the antigen into the lymphatic system. By this method one can avoid multiple ongoing injections and can also avoid the use of including professional antigen-presenting cells in the composition held in the reservoir.

The method of this invention can be viewed as inducing CTL immune response by providing high continuous local concentrations of antigen, which otherwise is quickly removed and degraded from the body after bolus injection. Potent activation of CD8+T cells requires signaling through the T cell receptor (TCR) in a manner that is dependent on both quantitative and qualitative factors. Quantitative factors refer to the number of TCRs engaged by peptide-MHC complexes. Qualitative considerations include the duration of engagement of the TCR by peptide-MHC complexes, with specific peptide-MHC complexes. Sustained regular deliveries of antigen allows optimal conditions to be established for inducing CD8+ T cells.

The antigen is delivered to the animal so that the antigen is present in the animal's lymphatic system on a sustained basis over a period of time. That is to say, it is delivered in such a way that the presence of the antigen is maintained over the period of time in the animal's lymphatic system. Thus, the antigen is delivered to the animal on a regular basis, i.e. the antigen is delivered regularly without significant interruption over the period of time. This regular delivery is achieved by the constant delivery of the antigen at low levels directly to the lymphatic system using an external device or an implantable device, as discussed hereinafter. Alternatively, the antigen can be delivered at higher levels to the animal by subcutaneous injection with indirect absorption or equilibration with the lymph system. Delivery on a regular basis is meant to include intermittent (stopping and transmitting at intervals) as well as continuous (transmitting without interruption) delivery. In intermittent delivery, the

times transmission is stopped will not be enough to reduce the level of antigen in the animal's lymphatic system to eliminate the desired specific immunological response. Thus, the antigen may be delivered in pulses or small doses over time.

Preferably, the sustained delivery is achieved by the positioning of a means of delivery so that the animal being treated does not have to receive multiple injections of the antigen, but instead has only one insertion of the means for delivery, e.g. an insertion of a catheter or needle for infusion of a suitable antigen-containing composition or the surgical implantation of an implantable device that release an appropriate, antigen-containing composition on a sustained basis.

The period of time over which the antigen will be released will be a time sufficient to induce and maintain the desired specific immunological response, e.g. to maintain a CTL response, and in the case of an animal with a tumor or infection, at a level sufficient to stimulate the immune system to attack the tumor and inhibit its growth or to attack the infection. Generally, this period of time may vary from a few days, e.g. a week, to a year or more. Preferably, the treatment, i.e. sustained delivery of the antigen, will extend for at least seven days and no more than six months. It has been found that the CTL response is induced by administration for at least seven days. To determine the period of time, the attending physician will evaluate, i.e., the severity of the condition, the strength of the patient, the antigenic response (e.g., the level of CD8+ cells measurable in the patient's system), the presence of toxic effects, and other factors known to one of skill in the art. Ultimately the time for sustained delivery in a cancer patient will be that necessary for improvement in the patient as evidenced by reduction in the size of the tumor, the rate of growth of the tumor, and/or the improvement in the overall health of the patient being treated. In the treatment of infectious diseases the treatment is continued until the health of the patient improves sufficiently to stop treatment.

The underlying immunological rationale for the utility of this invention arises from certain immunological considerations. The immune system has evolved to protect the host from microbial infection. CD4+ T cells together with B cells are the main components of the immune system humoral effector arm, which is crucial to eliminate extracellular pathogens or toxins. In contrast, the CD8+ T cell arm of the immune system is mainly responsible for eliminating intracellular pathogens, i.e. most importantly viruses, either via cytokine release or by cytotoxic activity. It is now emerging that these most efficient "killer cells" of the immune system would best



serve as the primary effector cells in tumor immunotherapy. An object of this invention is to mount a disease-specific CTL response (CD8+ T cell response) against the disease and sustain it over time, *e.g.*, a tumor specific or microbial specific CTL response.

5 CD8+ T cells recognize antigenic oligopeptides presented on HLA class I molecules of target cells, *e.g.*, tumor cells. The sequences of many HLA-A1 and HLA-A2 presented tumor and pathogen specific antigen peptides have recently been characterized. These peptides may be used in this invention to induce, *e.g.*, a melanoma-specific CD8+ T cell response. These peptides are discussed hereinafter.

10 In contrast to viral infection, class I-binding oligopeptides show only low immunogenicity. Most viruses induce peak CD8+ T cell responses around 7-10 days after systemic spread. This invention aims at enhancing the immunogenicity of class I binding oligopeptides by sustained, regular release of peptide into a lymphatic system and continued release into the lymphatic system.

15 In contrast to antibody-mediated B cell memory, which is long lived, T cell memory appears to be short lived or non-existent. In accordance with this invention, maintenance of functional T cell memory depends on persistence of antigen through continued, regular administration of the desired antigen. Having made this invention and looking at past concepts that might support this underlying rationale, some evidence includes the observation that delayed type hypersensitivity (DTH) of the tuberculin type (the only functional test for T cell memory in humans), can be elicited only in granulomatous disease, such as tuberculosis (tuberculin test), leprosy (lepromin test), brucellosis (brucellin test), sarcoidosis (Kveim test), Histoplasmosis (histoplasmin test) etc., but no such test could be established for non-granulomatous  
25 infectious disease. A factor that all granulomatous diseases have in common, is that the antigen persists within the granuloma - professional antigen presenting cells can use this reservoir to continuously restimulate specific T cells in lymphoid organs. In mice models (see Example 3) it is demonstrated that maintenance of functional CD8+ T cell memory was strictly dependent on continuous antigenic restimulation.

30 To determine whether a CTL response is obtained in an animal being treated in accordance with this invention, one measures the level of CD8+ cells (*i.e.* CTL) present in the blood or lymphatic organs such as the spleen or lymph nodes. This determination is done by first measuring the level of CD8+ cells before performing the method of this invention and measuring the level during treatment, *e.g.* at 7, 10,

20, 40 days, etc. The level or strength of the CD8<sup>+</sup> (CTL) response can be assessed *in vivo* or *in vitro*. In humans, there exists so far only one *in vivo* test to measure CD8<sup>+</sup> T cell responses, which is a skin test. In this skin test, HLA class I binding peptides are injected intradermally (such as described in Jäger, E. *et al.* Granulocyte-macrophage-colony-stimulating Factor Enhances Immune Responses To Melanoma-associated Peptides *in vivo* *Int. J. Cancer* 67, 54-62 (1996)). If a CTL response is present, these cells will recognize and attack peptide pulsed dermal cells, causing a local inflammatory reaction either via cytokine release or the cytotoxic mechanism (Kündig, T.M., Althage, A., Hengartner, H. & Zinkernagel, R.M. A skin test to assess CD8<sup>+</sup> cytotoxic T cell activity. *Proc. Natl. Acad. Sci. USA* 89:7757-776 (1992)). This inflammatory reaction can be quantified by measuring the diameter of the local skin rash and/or by measuring the diameter of the infiltrate (i.e., the swelling reaction). As an alternative to the injection of soluble free peptide, the HLA-class I binding peptide can also be injected intradermally in a bound form, e.g., bound to extracorporally derived dendritic cells. In other mammals, additional, although experimental, *in vivo* tests to assess CD8<sup>+</sup> T cell responses exist. For example, in a mouse model, CD8<sup>+</sup> T cell responses can be measured by challenge infection with a vaccinia recombinant virus expressing the peptide used for immunization. While naïve mice succumb to the infection with the vaccinia recombinant virus, mice with preexisting CD8<sup>+</sup> T cell immunity against the peptide epitope expressed by the vaccinia recombinant virus, are immune to reinfection. The level of immunity to reinfection can be quantified as the factor of reduction of the vaccinia virus titer recovered from mouse organs after challenge infection (Bachmann, M.F. & Kundig, T.M. *In vitro* vs. *in vivo* assays for the assessment of T- and B- cell function. *Curr. Opin. Immunol.* 6, 320-326 (1994)). For example, 5 days after challenge infection, a typical vaccinia recombinant virus titer recovered from a mouse ovary would be around 10<sup>7</sup> pfu per ovary, whereas the vaccinia recombinant virus titer in a mouse with a preexisting CD8<sup>+</sup> T cell response against the recombinant gene product would for example be around 10<sup>3</sup> pfu per ovary. Such a 10,000 fold-reduction in virus titer reflects biologically significant preexisting CD8<sup>+</sup> T cell activity against the recombinant gene product.

The level of CD8<sup>+</sup> T cell responses can also be quantified *in vitro*, by estimating the number of CD8<sup>+</sup> T cells specific for the antigenic peptide in question. In a naïve mammal the so called "frequency", i.e., the number of specific CD8<sup>+</sup> T

cells divided by the number of non-specific white blood cells, is less than  $10^{-6}$ . After successful immunization, the frequency increases due to proliferation of specific T cells. During an acute viral infection, for example, the frequency of specific CD8+ T cells may rise to  $10^{-2}$ . Then, after elimination of the virus, the frequency of specific CD8+ T cells usually drops to a "memory" level of around  $10^{-4}$ . Thus, the CD8+ T cell response can be quantified by measuring the frequency of specific CD8+ T cells. The higher the frequency, the stronger the response. The classical assays used to measure the frequency of specific CD8+ T cells are based on limiting dilution cell culture techniques, as described in detail by Kündig, T.M. et al. (On the role of antigen in maintaining cytotoxic T cell memory. *Proceedings of the National Academy of Sciences of the United States of America* 93, 9716-9723 (1996)). A novel approach to estimate the frequency of specific CD8+ T cells is to construct soluble class I MHC (for use in mice) or HLA molecules (for use in humans) with a peptide bound to their groove, so that the specific T cell receptors will bind to these complexes. These complexes can be labeled for detection, for example, with a fluorescent substance, allowing for detection by flow cytometry.

One current procedure to render peptides immunogenic is to inject them in context with "nature's most potent adjuvant", i.e., professional antigen presenting cells (APCs) such as dendritic cells (DCs) (Steinmann, R.M., The dendritic cells system and its role in immunogenicity, *Annual Review of Immunology* 9, 271-96 (1991)). DCs are the most potent APCs of the immune system. They can now be cultured in vitro by adding granulocyte macrophage colony stimulating factor (GM-CSF) and tumor necrosis factor alpha (TNF-alpha) or interleukin-4 (IL-4) to progenitors isolated from the blood of patients or mice (Inaba, K. et al., Identification of proliferating dendritic cell precursors in mouse blood, *Journal of Experimental Medicine* 175, 1157-1167 (1992)). Large numbers of DCs can then be pulsed with tumor specific antigen peptides and are injected back into the patient, where they migrate into lymphatic organs to induce T cell responses (Young, J.W. & Inaba, K., Dendritic Cells As Adjuvants For Class I Major Histocompatibility Complex-restricted Anti-tumor Immunity, *Journal of Experimental Medicine* 183, 7-11 (1996)). An object of this invention is to circumvent the time-consuming, labor intensive procedure of culturing DCs after isolation of DC progenitors and deliver the antigen to the lymphatic system free of APCs such as DCs. The method of this invention, i.e., the sustained, regular delivery of antigen into a lymphatic organ, allows sufficiently

high local concentrations of antigen inside the lymphatic organ, such that professional antigen presenting cells, e.g., dendritic cells, can be loaded with peptide *in vivo*. This can be viewed as a method of loading antigen presenting cells (dendritic cells) *in vivo* for inducing a CTL response.

5           The method of the present invention is clearly advantageous over the prior art methods for inducing a CTL response against a tumor or virus. For example, the present invention does not require repetitive immunizations to effect for prolonged anti-tumor immunotherapy. The sustained delivery of the antigen maintains the CTL response that could ultimately afford a prolonged aggressive posture of CTL against  
10 tumor cells, more thorough eradication, and protection against recurrence during the vaccine treatment. In the absence of antigen, CTL that have undergone primary activation soon cease to recirculate through the body, soon finding their way to the spleen where they become quiescent. Since CTL must immediately deliver a lethal hit, their residence in the spleen precludes an active role in protection against  
15 infections or tumor growth at distant sites in the body. The controlled release of antigen recognized by CTL in this invention circumvents this outcome as antigen delivery is maintained. Sustained released antigen delivery to the lymphatic system by this invention solves two major problems: it provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation that is  
20 necessary to keep CTL active, cytotoxic and recirculating through the body.

          Another fundamental improvement of the present method over prior art is that it facilitates the use of inherently non-immunogenic peptide antigens for CTL stimulation without the combined use of conventional adjuvants. This is very beneficial as most experimental adjuvants are toxic and poorly suited for use in  
25 humans. In addition adjuvants stimulate the TH2-type humoral immune response that negatively affects the CTL response. Further, since conventional adjuvants are not required, only the minimal antigenic epitope for a CTL response is required in the formulation.

          An additional advantage to the method of the present invention, where it  
30 embodies the use of mechanical delivery systems, is that the antigen delivery can be stopped if any adverse immunological effects are observed. For example, in vaccines against melanoma, CTL have been induced to attack not only malignant melanocytes but also healthy tissue, causing "vitiligo." The ability to discontinue a CTL vaccine at any time is a significant advance in vaccine safety. Peptides have a short half-life due

to catabolism in the liver. Therefore, the stimulation-effect falls soon after cessation of delivery.

As pointed out before, the method of this invention has two parts: (1) inducing an increased CTL response and (2) maintaining the response. The inducing and maintaining may be performed using the same device, as discussed hereinafter, or the inducing may be done separately, *e.g.*, by a separate injection of an antigen then following up with sustained delivery of the antigen over time to maintain the response.

#### Diseases treated according to the invention

In general, this invention is useful for treating an animal having (or being predisposed to) any disease to which the animal's immune system mounts a cell-mediated response to a disease-related antigen in order to attack the disease. Thus, the type of disease may be a malignant tumor or a chronic infectious disease caused by a bacterium, virus, protozoan, helminth, or other microbial pathogen that enters intracellularly and is attacked, *i.e.*, by the cytotoxic T lymphocytes. In addition, the invention is useful for treating an animal that may be at risk of developing such diseases.

#### Malignant Tumors

In a mature animal, a balance usually is maintained between cell renewal and cell death in most organs and tissues. The various types of mature cells in the body have a given life span; as these cells die, new cells are generated by the proliferation and differentiation of various types of stem cells. Under normal circumstances, the production of new cells is so regulated that the numbers of any particular type of cell remain constant. Occasionally, though, cells arise that are no longer responsive to normal growth-control mechanisms. These cells give rise to clones of cells that can expand to a considerable size, producing a *tumor*, or *neoplasm*. A tumor that is not capable of indefinite growth and does not invade the healthy surrounding tissue extensively is *benign*. A tumor that continues to grow and becomes progressively invasive is *malignant*; the term *cancer* refers specifically to a malignant tumor. In addition to uncontrolled growth, malignant tumors exhibit *metastasis*; in this process, small clusters of cancerous cells dislodge from a tumor, invade the blood or lymphatic vessels, and are carried to other tissues, where they continue to proliferate. In this way a primary tumor at one site can give rise to a secondary tumor at another site.

The methods, devices and articles of manufacture discussed herein are useful for treating animals having malignant tumors.

Malignant tumors treated according to this invention are classified according to the embryonic origin of the tissue from which the tumor is derived. *Carcinomas* are tumors arising from endodermal or ectodermal tissues such as skin or the epithelial lining of internal organs and glands. A melanoma is a type of carcinoma of the skin for which this invention is particularly useful. *Sarcomas*, which arise less frequently, are derived from mesodermal connective tissues such as bone, fat, and cartilage. The *leukemias* and *lymphomas* are malignant tumors of hematopoietic cells of the bone marrow. Leukemias proliferate as single cells, whereas lymphomas tend to grow as tumor masses. The malignant tumors may show up at numerous organs or tissues of the body to establish a cancer. The types of cancer that can be treated in accordance with this invention include the following: bladder, brain, breast, cervical, colo-rectal, esophageal, kidney, liver, lung, nasopharyngeal, pancreatic, prostate, skin, stomach, uterine, and the like. The present invention is not limited to the treatment of an existing tumor or infectious disease but can also be used to prevent or lower the risk of developing such diseases in an individual, *i.e.*, for prophylactic use. Potential candidates for prophylactic vaccination include individuals with a high risk of developing cancer, *i.e.*, with a personal or tuminal history of certain types of cancer.

The incidence of skin cancer has increased substantially over the last decades. Lifetime analysis indicates that around 1/1500 humans born in 1935, 1/600 born in 1960, 1/100 born in 1990 and a projected 1/75 humans born in the year 2000 will have melanoma in their lifetime. Surgical excision usually cures melanoma. However, even small looking lesions may have already metastasized at the time of diagnosis. The prognosis of metastasized melanoma is very poor and correlates with the thickness of the primary tumor and with its localization.

The current treatment of malignant melanoma aims at surgical removal of the primary tumor. If metastases are present, chemotherapy and biological response modifiers are additionally used. However, patients with stage IV malignant melanoma are almost invariably incurable and treatments are palliative. Patients with Stage IV malignant melanoma have a median survival time of approximately one year and only a 10% chance of long-term survival. There is at present no generally accepted standard therapy for metastatic melanoma. Objective response rates to mono- or polychemotherapy are low in comparison with other tumors, reaching no

more than 15-35%. An improved treatment outcome in stage IV malignant melanoma seems unachievable either by chemotherapeutic combinations or by increasing doses to levels where autologous bone marrow transplantation becomes necessary. The method of this invention is useful for treating malignant melanoma, even at Stage IV.

## 5 **Infectious Diseases**

Infectious diseases, which have plagued animal populations (particularly humans) throughout history, still cause millions of deaths each year. The infectious diseases that can be treated using this invention include those caused by pathogens such as bacteria, viruses, protozoa, helminths, and the like. These diseases include  
10 such chronic diseases such as acute respiratory infections, diarrheal diseases, tuberculosis, malaria, hepatitis (hepatitis A, B C, D, E, F virus), measles, mononucleosis (Epstein-Barr virus), whooping cough (pertussis), AIDS (human immunodeficiency virus 1 & 2), rabies, yellow fever, and the like. Other diseases caused by human papilloma virus or various strains of virus are treatable by this  
15 method.

In some instances, the mammal, in particular human, can be treated prophylactically, such as when there may be a risk of developing disease. An individual travelling to or living in an area of endemic infectious disease may be considered to be at risk and a candidate for prophylactic vaccination against the  
20 particular infectious agent. For example, the CTL response can be induced in a human expecting to enter a malarial area and/or while in the malarial area by using a CTL-inducing, malaria-specific antigen to lower the risk of developing malaria. Preventative treatment can be applied to any number of diseases including those listed above, where there is a known relationship between the particular disease and a  
25 particular risk factor, such as geographical location or work environment.

## **Antigens useful in the invention**

An antigen useful in this invention is one that stimulates the immune system of a mammal having a malignant tumor or infectious disease to attack the tumor and inhibit its growth or to destroy the pathogen causing the disease. Thus, the antigen  
30 used in the invention is matched to the specific disease found in the animal being treated. In this regard the antigen may be said to induce a CTL response (also referred to as a cell-mediated immune response), i.e. a cytotoxic reaction by the immune system that results in lysis of the target cells (e.g., the malignant tumor cells or pathogen-infected cells).

To determine whether an antigen is matched to a particular patient, whether human or other animal, the tissue type of the patient is first determined. If human, the tissue must demonstrate the appropriate human leukocyte antigen (HLA) capable of binding and displaying the antigen to CTL. It is preferable that the HLA typing be performed on the target cells, since a significant portion of tumors escape immune detection by downregulating expression of HLA. Therefore HLA expression on normal cells of the patient does not necessarily reflect that found on tumor cells in their body. A tumor from a patient is also screened to determine if he or she expresses the antigen that is being used in the vaccine formulation.

Immunohistochemistry and/or polymerase chain reaction (PCR) techniques both can be used to detect antigen in the tumor cells. Immunohistochemistry offers the advantage in that it stains a cross-section of tumor in a slide preparation, allowing investigators to observe the antigen expression pattern in cross-section of tumor, which is typically heterogeneous for antigen expression. PCR has the advantage of not requiring specific monoclonal antibodies for staining and is a fast and powerful technique. In addition, PCR can be applied in situ. Ideally, both immunohistochemical and PCR methods should be combined when assessing antigen expression in tumors. While the antigen compositions useful in this invention are designed to include the most commonly expressed tumor antigens (as discussed hereafter), not all tumors will express the desired antigen(s). Where a tumor fails to express the desired antigen, the patient is excluded for consideration for that particular antigen composition. Thus, an aspect of this invention is a process for preparing a device useful for providing a sustained CTL response over time by matching a subject's antigen specific to the tumor or pathogen in the subject, preparing a physiologically-acceptable composition of the antigen so matched, and combining the composition in a suitable delivery device as discussed in hereinafter.

Immune activation of CD8+ T cells generates a population of effector cells with lytic capability called cytotoxic T lymphocytes, or CTL. These effector cells have important roles in the recognition and elimination of malignant cells and pathogens. In general, CTL are CD8+ and are therefore class I MHC restricted, although in rare instances CD4+ class II - restricted T cells have been shown to function as CTL. Since virtually all nucleated cells in the body express class I MHC molecules, CTL can recognize and eliminate almost any altered body cell. CD8+T



cells recognize antigen presented on HLA class I molecules of tumor cells through T cell receptors.

The CTL-mediated immune response can be divided into two phases, reflecting different aspects of the cytotoxic T-cell response. The first phase involves the activation and differentiation of T<sub>c</sub> (CD8+) cells into functional effector CTLs. In the second phase, CTLs recognize antigen - class I MHC complexes on specific target cells, initiating a sequence of events that culminates in target-cell destruction. Further detailed discussion of the process is found at Chapter 15 of the Second Edition of "Immunology" by Janis Kuby, W.H. Freeman and Company (1991).

The type of tumor antigen useful in this invention may be a *tumor-specific antigen (TSA)* or a *tumor-associated antigen (TAA)*. A *TSA* is unique to tumor cells and does not occur on other cells in the body. A *TAA* associated antigen is not unique to a tumor cell and instead is also expressed on a normal cell under conditions that fail to induce a state of immunologic tolerance to the antigen. The expression of the antigen on the tumor may occur under conditions that enable the immune system to respond to the antigen. *TAA*s may be antigens that are expressed on normal cells during fetal development when the immune system is immature and unable to respond or they may be antigens that are normally present at extremely low levels on normal cells but which are expressed at much higher levels on tumor cells. *TSAs* and *TAA*s can be jointly referred to as *TSA* or a tumor related antigen.

Tumor antigens useful in the present invention, whether tumor-specific or tumor-associated, must be capable of inducing a CTL-mediated immune response. The presence of tumor antigens that elicit a cell-mediated response has been demonstrated by the rejection of tumors transplanted into syngeneic recipients; because of this phenomenon, these tumor antigens are referred to as *tumor-specific transplantation antigens (TSTAs)* or *tumor-associated transplantation antigens (TATAs)*. It has been difficult to characterize tumor transplantation antigens because they do not generally elicit an antibody response and therefore they cannot be isolated by immunoprecipitation. Many are peptides that are presented together with MHC molecules on the surface of tumor cells and have been characterized by their ability to induce an antigen-specific CTL.

The type of pathogen specific antigen useful in this invention may be short oligopeptides derived from pathogen proteins. These oligopeptides must bind to class I MHC (for use in mice), class I HLA (for use in humans), or class I molecules of any

other mammals. Also, such class I molecule bound peptides should be recognizable by specific T cell receptors. Such oligopeptides usually have a length of 8-15 amino acids. Several examples of such pathogen derived oligopeptides, so called T cell epitopes, are given in Tables I and II.

5       The tumor antigens and pathogen-specific antigens useful in this invention are generally thought to be presented at the surface of an antigen presenting cell (APC) to stimulate the immune system through class I molecules of the major histocompatibility complex (MHC) interactively with the CD8+ cells.

10       Antigens useful in the invention are generally protein-based entities of a molecular weight of up to 100,000 daltons. Appropriate antigens include, but are not limited to differentiation antigens, tumor-specific multilineage antigens, embryonic antigens, antigens of oncogenes and mutated tumor-suppressor genes, unique tumor antigens resulting from chromosomal translocations, viral antigens, and others that may be apparent presently or in the future to one of skill in the art. It is preferable  
15       that the antigen be a peptide of 8 to 15 amino acids in length that is an epitope of a larger antigen, i.e. it is a peptide having an amino acid sequence corresponding to the site on the larger molecule that is recognized and bound by a particular T-cell receptor. These smaller peptides are available to one of skill in the art by following the teachings of U.S. Patents 5,747,269 to Rammensee et al. issued May 5, 1998;  
20       5,698,396 to Pfreundschuh issued December 16, 1997; and PCT Application Numbers PCT/EP95/02593 filed 4 July 1995, PCT/DE96/00351 filed 26 Feb 1996, and PCT/EP97/05198 filed 22 Sept 1997, all of which are incorporated herein by reference.

25       A powerful method has been recently developed for identifying new peptides that are useful in the invention. Genes determined to express protein with high exclusivity in tumor cells or microbial cells (e.g. viruses) can be identified using a so called SEREX process, which involves expression cloning using tumor cell libraries and screening these libraries against immunoglobulin in patient sera. Over one hundred genes have recently been identified from tumor biopsies using this process.  
30       These genes can now be used in a peptide prediction algorithm developed by Hans-Georg Rammensee. Algorithms have been developed for all major HLA types found in the human population. First the protein sequence is "translated" based on the gene sequence. The algorithms can predict peptide epitopes for various HLA types based on the protein sequence. Since the predicted peptides are indeed predictions and are

not always naturally found on cells, tumor samples are used to confirm the predicted peptides by actually isolating minute trace peptide from tumors. Being able to calculate the exact mass of the predicted peptides allows trace peptide identification using ultrasensitive mass spectrophotometry, which can detect peptides in quantities less than that which would permit peptide sequencing and identification. Once these tumor-associated peptides have been identified they are suitable for use in the invention, since peptides of a known sequence may be synthesized in large quantities (several grams) providing for sufficient amounts of peptides for use in this invention.

Thus it can be seen that another aspect of this invention is a process for preparing a composition useful in a device of this invention as discussed hereinafter. The process comprises identifying a gene determined to express a protein with high exclusivity in a tumor or microbial cell, cloning cell libraries; screening the libraries against immunoglobulin in patient sera, using the algorithm defined in the literature developed by Hans-George Rammensee to predict an epitope for the HLA type protein based on the gene sequence, matching the predicted antigen sequence to a patient tumor sample, isolating the matched antigen, and preparing a composition of the antigen for use in a delivery device as discussed hereinafter.

Examples of large, protein-based antigens include the following:

differentiation antigens such as MART-1/MelanA (MART-I), gp100 (Pmel 17), tyrosinase, TRP-1, TRP-2 and tumor-specific multilineage antigens such as MAGE-1, MAGE-3, BAGE, GAGE-1, GAGE-2, p15; overexpressed embryonic antigens such as CEA; overexpressed oncogenes and mutated tumor-suppressor genes such as p53, Ras, HER-2/neu; unique tumor antigens resulting from chromosomal translocations such as BCR-ABL, E2A-PRL, H4-RET, IGH-IGK, MYL-RAR; and viral antigens, such as the Epstein Barr virus antigens EBVA and the human papillomavirus (HPV) antigens E6 and E7. Other large, protein-based antigens include TSP-180, MAGE-4, MAGE-5, MAGE-6, RAGE, NY-ESO, p185erbB2, p180erbB-3, c-met, nm-23H1, PSA, TAG-72, CA 19-9, CA 72-4, CAM 17.1, NuMa, K-ras,  $\beta$ -Catenin, CDK4, Mum-1, p15, p16. These protein-based antigens are known and available to those of skill in the art in the literature or commercially.

Examples of peptide antigens of 8-15 amino acids include those set forth in Table I, Table II, and Table III. Table I sets forth antigens that are virally derived. The Table shows the virus type, the protein expressed by the virus, the amino acid (AA) position on the viral protein, the AA sequence of the T-cell epitope/MHC

ligand, the type of MHC molecule presenting the antigen, and a reference source. A more complete list is provided in the book by Han-Georg Rammensee, Jutta Bachmann, and Stefan Stevanovic entitled "MHC Ligands and Peptide Motifs," Springer-Verlag, Germany, 1997 Landes Bioscience, Austin, Texas). The reference number given in Table I is the same number (and reference source) given in Table 5.3 of the above Rammensee book, all of which is incorporated herein by reference.

Table I: Viral epitopes on MHC class 1 molecules

| Virus          | Protein        | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule       | Ref.     |
|----------------|----------------|-------------|--------------------------------------|--------------------|----------|
| Adenovirus 3   | E3 9Kd         | 30-38       | LIVIGILIL<br>(SEQ. ID NO.:1)         | HLA-A*0201         | 104      |
| Adenovirus 5   | E1A            | 234-243     | SGPSNTPPEI<br>(SEQ. ID NO.:2)        | H2-D <sup>b</sup>  | 105      |
| Adenovirus 5   | E1B            | 192-200     | VNIRNCCYI<br>(SEQ. ID NO.:3)         | H2-D <sup>b</sup>  | 106      |
| Adenovirus 5   | E1A            | 234-243     | SGPSNIPPEI (T>I)<br>(SEQ. ID NO.:4)  | H2-D <sup>b</sup>  | 106      |
| CSFV           | NS polyprotein | 2276-2284   | ENALLVALF<br>(SEQ. ID NO.:5)         | SLA, haplotype d/d | 107      |
| Dengue virus 4 | NS3            | 500-508     | TPEGIITL<br>(SEQ. ID NO.:6)          | HLA-B*3501         | 108, 109 |
| EBV            | LMP-2          | 426-434     | CLGGLTLMV<br>(SEQ. ID NO.:7)         | HLA-A*0201         | 110      |
| EBV            | EBNA-1         | 480-484     | NIAEGLRAL<br>(SEQ. ID NO.:8)         | HLA-A*0201         | 111      |
| EBV            | EBNA-1         | 519-527     | NLRRGTALA<br>(SEQ. ID NO.:9)         | HLA-A*0201         | 111      |
| EBV            | EBNA-1         | 525-533     | ALAIPQCRL<br>(SEQ. ID NO.:10)        | HLA-A*0201         | 111      |
| EBV            | EBNA-1         | 575-582     | VLKDAIKDL<br>(SEQ. ID NO.:11)        | HLA-A*0201         | 111      |
| EBV            | EBNA-1         | 562-570     | FMVFLQTHI<br>(SEQ. ID NO.:12)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 15-23       | HLIVDTDSL<br>(SEQ. ID NO.:13)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 22-30       | SLGNPSLSV<br>(SEQ. ID NO.:14)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 126-134     | PLASAMRML<br>(SEQ. ID NO.:15)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 132-140     | RMLWMANYI<br>(SEQ. ID NO.:16)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 133-141     | MLWMANYIV<br>(SEQ. ID NO.:17)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 151-159     | ILPQGPQTA<br>(SEQ. ID NO.:18)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 171-179     | PLRPTAPTI<br>(SEQ. ID NO.:19)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 205-213     | PLPPATLTV<br>(SEQ. ID NO.:20)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 246-254     | RMHLPVLHV<br>(SEQ. ID NO.:21)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 287-295     | PMPLPPSQL<br>(SEQ. ID NO.:22)        | HLA-A*0201         | 111      |
| EBV            | EBNA-2         | 294-302     | QLPPPAAPA<br>(SEQ. ID NO.:23)        | HLA-A*0201         | 111      |

Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule | Ref.     |
|-------|---------|-------------|--------------------------------------|--------------|----------|
| EBV   | EBNA-2  | 381-389     | SMPELSPVL<br>(SEQ. ID NO.:24)        | HLA-A*0201   | 111      |
| EBV   | EBNA-2  | 453-461     | DLDESWDYI<br>(SEQ. ID NO.:25)        | HLA-A*0201   | 111      |
| EBV   | BZLF1   | 43-51       | PLPCVLWPV<br>(SEQ. ID NO.:26)        | HLA-A*0201   | 111      |
| EBV   | BZLF1   | 167-175     | SLEECDSEL<br>(SEQ. ID NO.:27)        | HLA-A*0201   | 111      |
| EBV   | BZLF1   | 176-184     | EIKRYKNRV<br>(SEQ. ID NO.:28)        | HLA-A*0201   | 111      |
| EBV   | BZLF1   | 195-203     | QLLQHYREV<br>(SEQ. ID NO.:29)        | HLA-A*0201   | 111      |
| EBV   | BZLF1   | 196-204     | LLQHYREVA<br>(SEQ. ID NO.:30)        | HLA-A*0201   | 111      |
| EBV   | BZLF1   | 217-225     | LLKQMCPSL<br>(SEQ. ID NO.:31)        | HLA-A*0201   | 111      |
| EBV   | BZLF1   | 229-237     | SIIPRTPDV<br>(SEQ. ID NO.:32)        | HLA-A*0201   | 111      |
| EBV   | EBNA-6  | 284-293     | LLDFVRFMGV<br>(SEQ. ID NO.:33)       | HLA-A*0201   | 112      |
| EBV   | EBNA-3  | 464-472     | SVRDLRLARL<br>(SEQ. ID NO.:34)       | HLA-A*0203   | 113      |
| EBV   | EBNA-4  | 416-424     | IVTDFSVIK<br>(SEQ. ID NO.:35)        | HLA-A*1101   | 114, 115 |
| EBV   | EBNA-4  | 399-408     | AVFDRKSDAK<br>(SEQ. ID NO.:36)       | HLA-A*0201   | 116      |
| EBV   | EBNA-3  | 246-253     | RYSIFFDY<br>(SEQ. ID NO.:37)         | HLA-A24      | 113      |
| EBV   | EBNA-6  | 881-889     | QPRAPIRPI<br>(SEQ. ID NO.:38)        | HLA-B7       | 117      |
| EBV   | EBNA-3  | 379-387     | RPPIFIRRL<br>(SEQ. ID NO.:39)        | HLA-B7       | 117      |
| EBV   | EBNA-1  | 426-434     | EPDVPPGAI<br>(SEQ. ID NO.:40)        | HLA-B7       | 111      |
| EBV   | EBNA-1  | 228-236     | IPQCRLTPL<br>(SEQ. ID NO.:41)        | HLA-B7       | 111      |
| EBV   | EBNA-1  | 546-554     | GPGPQPGPL<br>(SEQ. ID NO.:42)        | HLA-B7       | 111      |
| EBV   | EBNA-1  | 550-558     | QPGPLRESI<br>(SEQ. ID NO.:43)        | HLA-B7       | 111      |
| EBV   | EBNA-1  | 72-80       | RPQKRPSCI<br>(SEQ. ID NO.:44)        | HLA-B7       | 111      |
| EBV   | EBNA-2  | 224-232     | PPTPLLTVL<br>(SEQ. ID NO.:45)        | HLA-B7       | 111      |
| EBV   | EBNA-2  | 241-249     | TPSPPRMHL<br>(SEQ. ID NO.:46)        | HLA-B7       | 111      |
| EBV   | EBNA-2  | 244-252     | PPRMHLPVL<br>(SEQ. ID NO.:47)        | HLA-B7       | 111      |

Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein      | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref. |
|-------|--------------|-------------|--------------------------------------|-------------------|------|
| EBV   | EBNA-2       | 254-262     | VPDQSMHPL<br>(SEQ. ID NO.:48)        | HLA-B7            | 111  |
| EBV   | EBNA-2       | 446-454     | PPSIDPADL<br>(SEQ. ID NO.:49)        | HLA-B7            | 111  |
| EBV   | BZLF1        | 44-52       | LPCVLWPVL<br>(SEQ. ID NO.:50)        | HLA-B7            | 111  |
| EBV   | BZLF1        | 222-231     | CPSLDVDSII<br>(SEQ. ID NO.:51)       | HLA-B7            | 111  |
| EBV   | BZLF1        | 234-242     | TPDVLHEDL<br>(SEQ. ID NO.:52)        | HLA-B7            | 111  |
| EBV   | EBNA-3       | 339-347     | FLRGRAYGL<br>(SEQ. ID NO.:53)        | HLA-B8            | 118  |
| EBV   | EBNA-3       | 26-34       | QAKWRLQTL<br>(SEQ. ID NO.:54)        | HLA-B8            | 113  |
| EBV   | EBNA-3       | 325-333     | AYPLHEQHG<br>(SEQ. ID NO.:55)        | HLA-B8            | 116  |
| EBV   | EBNA-3       | 158-166     | YIKSFVSDA<br>(SEQ. ID NO.:56)        | HLA-B8            | 116  |
| EBV   | LMP-2        | 236-244     | RRRWRLTV<br>(SEQ. ID NO.:57)         | HLA-B*2704        | 119  |
| EBV   | EBNA-6       | 258-266     | RRIYDLIEL<br>(SEQ. ID NO.:58)        | HLA-B*2705        | 119  |
| EBV   | EBNA-3       | 458-466     | YPLHEQHGM<br>(SEQ. ID NO.:59)        | HLA-B*3501        | 120  |
| EBV   | EBNA-3       | 458-466     | YPLHEQHGM<br>(SEQ. ID NO.:59)        | HLA-B*3503        | 113  |
| HCV   | NS3          | 389-397     | HSKKKCDEL<br>(SEQ. ID NO.:60)        | HLA-B8            | 145  |
| HCV   | env E        | 44-51       | ASRCWVAM<br>(SEQ. ID NO.:61)         | HLA-B*3501        | 146  |
| HCV   | core protein | 27-35       | GQIVGGVYL<br>(SEQ. ID NO.:62)        | HLA-B*40012       | 147  |
| HCV   | NS1          | 77-85       | RPLTDFDQGW<br>(SEQ. ID NO.:63)       | HLA-B*5301        | 145  |
| HCV   | core protein | 18-27       | LMGYIPLVGA<br>(SEQ. ID NO.:64)       | H2-D <sup>d</sup> | 138  |
| HCV   | core protein | 16-25       | ADLMGYIPLV<br>(SEQ. ID NO.:65)       | H2-D <sup>d</sup> | 148  |
| HCV   | NS5          | 409-424     | MSYSWTGALVTPCAEE<br>(SEQ. ID NO.:66) | H2-D <sup>d</sup> | 149  |
| HCV   | NS1          | 205-213     | KHPDATYSR<br>(SEQ. ID NO.:67)        | Papa-A06          | 150  |
| HCV-1 | NS3          | 400-409     | KLVALGINAV<br>(SEQ. ID NO.:68)       | HLA-A*0201        | 141  |
| HCV-1 | NS3          | 440-448     | GDFDSVIDC<br>(SEQ. ID NO.:69)        | Patr-B16          | 151  |
| HCV-1 | env E        | 118-126     | GNASRCWVA<br>(SEQ. ID NO.:70)        | Patr-B16          | 151  |

Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref.     |
|-------|---------|-------------|--------------------------------------|-------------------|----------|
| HCV-1 | NS1     | 159-167     | TRPPLGNWF<br>(SEQ. ID NO.:71)        | Patr-B13          | 151      |
| HCV-1 | NS3     | 351-359     | VPHPNIEEV<br>(SEQ. ID NO.:72)        | Patr-B13          | 151      |
| HCV-1 | NS3     | 438-446     | YTGFDFDSVI<br>(SEQ. ID NO.:73)       | Patr-B01          | 151      |
| HCV-1 | NS1     | 328-335     | SWAIKWEY<br>(SEQ. ID NO.:74)         | Patr-A11          | 151      |
| HCV-1 | NS1     | 205-213     | KHPDATYSR<br>(SEQ. ID NO.:75)        | Patr-A04          | 150      |
| HCV-1 | NS3     | 440-448     | GDFDSVIDC<br>(SEQ. ID NO.:76)        | Patr-A04          | 150      |
| HIV   | gp41    | 583-591     | RYLKDQQLL<br>(SEQ. ID NO.:77)        | HLA-A24           | 152      |
| HIV   | gagp24  | 267-275     | IVGLNKIVR<br>(SEQ. ID NO.:78)        | HLA-A*3302        | 153, 154 |
| HIV   | gagp24  | 262-270     | EIYKRWIIL<br>(SEQ. ID NO.:79)        | HLA-B8            | 155, 156 |
| HIV   | gagp24  | 261-269     | GEIYKRWII<br>(SEQ. ID NO.:80)        | HLA-B8            | 155, 156 |
| HIV   | gagp17  | 93-101      | EIKDTKEAL<br>(SEQ. ID NO.:81)        | HLA-B8            | 155, 157 |
| HIV   | gp41    | 586-593     | YLKDQQLL<br>(SEQ. ID NO.:82)         | HLA-B8            | 158      |
| HIV   | gagp24  | 267-277     | ILGLNKIVRMY<br>(SEQ. ID NO.:83)      | HLA-B*1501        | 153      |
| HIV   | gp41    | 584-592     | ERYLKDQQL<br>(SEQ. ID NO.:84)        | HLA-B14           | 158      |
| HIV   | nef     | 115-125     | YHTQGYFPQWQ<br>(SEQ. ID NO.:85)      | HLA-B17           | 159      |
| HIV   | nef     | 117-128     | TQGYFPQWQNYT<br>(SEQ. ID NO.:86)     | HLA-B17           | 159      |
| HIV   | gp120   | 314-322     | GRAFVTIGK<br>(SEQ. ID NO.:87)        | HLA-B*2705        | 160, 184 |
| HIV   | gagp24  | 263-271     | KRWIILGLN<br>(SEQ. ID NO.:88)        | HLA-B*2702        | 161      |
| HIV   | nef     | 72-82       | QVPLRPMTYK<br>(SEQ. ID NO.:89)       | HLA-B*3501        | 159      |
| HIV   | nef     | 117-125     | TQGYFPQWQ<br>(SEQ. ID NO.:90)        | HLA-B*3701        | 159      |
| HIV   | gagp24  | 143-151     | HQAISPRTL<br>(SEQ. ID NO.:91)        | HLA-Cw*0301       | 162      |
| HIV   | gagp24  | 140-151     | QMVHQAI SPRTL<br>(SEQ. ID NO.:92)    | HLA-Cw*0301       | 162      |
| HIV   | gp120   | 431-440     | MYAPPIGGQI<br>(SEQ. ID NO.:93)       | H2-K <sup>d</sup> | 163      |
| HIV   | gp160   | 318-327     | RGPGRAFVTI<br>(SEQ. ID NO.:94)       | H2-D <sup>d</sup> | 164, 165 |



Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref.     |
|-------|---------|-------------|--------------------------------------|-------------------|----------|
| HIV   | gp120   | 17-29       | MPGRAFTI<br>(SEQ. ID NO.:95)         | H2-L <sup>a</sup> | 166, 167 |
| HIV-1 | RT      | 476-484     | ILKEPVHGV<br>(SEQ. ID NO.:96)        | HLA-A*0201        | 168, 169 |
| HIV-1 | nef     | 190-198     | AFHHVAREL<br>(SEQ. ID NO.:97)        | HLA-A*0201        | 170      |
| HIV-1 | gp160   | 120-128     | KLTPLCVTL<br>(SEQ. ID NO.:98)        | HLA-A*0201        | 171      |
| HIV-1 | gp160   | 814-823     | SLLNATDIAV<br>(SEQ. ID NO.:99)       | HLA-A*0201        | 171      |
| HIV-1 | RT      | 179-187     | VIYQYMDDL<br>(SEQ. ID NO.:100)       | HLA-A*0201        | 172      |
| HIV-1 | gagp17  | 77-85       | SLYNTVATL<br>(SEQ. ID NO.:101)       | HLA-A*0201        | 173      |
| HIV-1 | gp160   | 315-329     | RGPRAFTI<br>(SEQ. ID NO.:102)        | HLA-A*0201        | 174      |
| HIV-1 | gp11    | 768-778     | RLRDLLIVTR<br>(SEQ. ID NO.:103)      | HLA-A3            | 175, 178 |
| HIV-1 | nef     | 73-82       | QVPLRPMTYK<br>(SEQ. ID NO.:104)      | HLA-A3            | 176      |
| HIV-1 | gp120   | 36-45       | TVYYGVPVWK<br>(SEQ. ID NO.:105)      | HLA-A3            | 177      |
| HIV-1 | gagp17  | 20-29       | RLRPGGKKK<br>(SEQ. ID NO.:106)       | HLA-A3            | 177      |
| HIV-1 | gp120   | 38-46       | VYYGVPVWK<br>(SEQ. ID NO.:107)       | HLA-A3            | 179      |
| HIV-1 | nef     | 74-82       | VPLRPMTYK<br>(SEQ. ID NO.:108)       | HLA-a*1101        | 114      |
| HIV-1 | gagp24  | 325-333     | AIFQSSMTK<br>(SEQ. ID NO.:109)       | HLA-A*1101        | 114      |
| HIV-1 | nef     | 73-82       | QVPLRPMTYK<br>(SEQ. ID NO.:104)      | HLA-A*1101        | 180      |
| HIV-1 | nef     | 83-94       | AAVDLSHFLKEK<br>(SEQ. ID NO.:110)    | HLA-A*1101        | 159      |
| HIV-1 | gagp24  | 349-359     | ACQGVGGPGGHK<br>(SEQ. ID NO.:111)    | HLA-A*1101        | 181      |
| HIV-1 | gagp24  | 203-212     | ETINEEAAEW<br>(SEQ. ID NO.:112)      | HLA-A25           | 182      |
| HIV-1 | nef     | 128-137     | TPGPGVRYPL<br>(SEQ. ID NO.:113)      | HLA-B7            | 159      |
| HIV-1 | gagp17  | 24-31       | GGKKKYKL<br>(SEQ. ID NO.:114)        | HLA-B8            | 183      |
| HIV-1 | gp120   | 2-10        | RVKEKYQHL<br>(SEQ. ID NO.:115)       | HLA-B8            | 181      |
| HIV-1 | gagp24  | 298-306     | DRFYKTLRA<br>(SEQ. ID NO.:116)       | HLA-B14           | 173      |
| HIV-1 | NEF     | 132-147     | GVRYPITFGWCYKLP<br>(SEQ. ID NO.:117) | HLA-B18           | 159      |

Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule | Ref.          |
|-------|---------|-------------|--------------------------------------|--------------|---------------|
| HIV-1 | gagp24  | 265-24      | KRWIILGLNK<br>(SEQ. ID NO.:118)      | HLA-B*2705   | 184, 153      |
| HIV-1 | nef     | 190-198     | AFHHVAREL<br>(SEQ. ID NO.:97)        | HLA-B*5201   | 170           |
| EBV   | EBNA-6  | 335-343     | KEHVIQNAF<br>(SEQ. ID NO.:119)       | HLA-B44      | 121           |
| EBV   | EBNA-6  | 130-139     | EENLLDFVRF<br>(SEQ. ID NO.:120)      | HLA-B*4403   | 122           |
| EBV   | EBNA-2  | 42-51       | DTPLIPLTIF<br>(SEQ. ID NO.:121)      | HLA-B51      | 121           |
| EBV   | EBNA-6  | 213-222     | QNGALAINTF<br>(SEQ. ID NO.:122)      | HLA-B62      | 112           |
| EBV   | EBNA-3  | 603-611     | RLRAEAGVK<br>(SEQ. ID NO.:123)       | HLA-A3       | 123           |
| HBV   | sAg     | 348-357     | GLSPTVWLSV<br>(SEQ. ID NO.:124)      | HLA-A*0201   | 124           |
| HBV   | SAg     | 335-343     | WLSLLVPFV<br>(SEQ. ID NO.:125)       | HLA-A*0201   | 124           |
| HBV   | cAg     | 18-27       | FLPSDFFPSV<br>(SEQ. ID NO.:126)      | HLA-A*0201   | 125, 126, 127 |
| HBV   | cAg     | 18-27       | FLPSDFFPSV<br>(SEQ. ID NO.:126)      | HLA-A*0202   | 127           |
| HBV   | cAg     | 18-27       | FLPSDFFPSV<br>(SEQ. ID NO.:126)      | HLA-A*0205   | 127           |
| HBV   | cAg     | 18-27       | FLPSDFFPSV<br>(SEQ. ID NO.:126)      | HLA-A*0206   | 127           |
| HBV   | pol     | 575-583     | FLLSLGIHL<br>(SEQ. ID NO.:127)       | HLA-A*0201   | 128           |
| HBV   | pol     | 816-824     | SLYADSPSV<br>(SEQ. ID NO.:128)       | HLA-A*0201   | 128           |
| HBV   | pol     | 455-463     | GLSRYVARL<br>(SEQ. ID NO.:129)       | HLA-A*0201   | 128           |
| HBV   | env     | 338-347     | LLVPFVQWFV<br>(SEQ. ID NO.:130)      | HLA-A*0201   | 129           |
| HBV   | pol     | 642-650     | ALMPYACI<br>(SEQ. ID NO.:131)        | HLA-A*0201   | 129           |
| HBV   | env     | 378-387     | LLPIFFCLWV<br>(SEQ. ID NO.:132)      | HLA-A*0201   | 129           |
| HBV   | pol     | 538-546     | YMDDVVLGA<br>(SEQ. ID NO.:133)       | HLA-A*0201   | 129           |
| HBV   | env     | 250-258     | LLLCLIFLL<br>(SEQ. ID NO.:134)       | HLA-A*0201   | 130           |
| HBV   | env     | 260-269     | LLDYQGMLPV<br>(SEQ. ID NO.:135)      | HLA-A*0201   | 130           |
| HBV   | env     | 370-379     | SIVSPFIPLL<br>(SEQ. ID NO.:136)      | HLA-A*0201   | 130           |
| HBV   | env     | 183-191     | FLLTRILT<br>(SEQ. ID NO.:137)        | HLA-A*0201   | 130           |

Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref. |
|-------|---------|-------------|--------------------------------------|-------------------|------|
| HBV   | cAg     | 88-96       | YVNVNMGLK<br>(SEQ. ID NO.:138)       | HLA-A*1101        | 131  |
| HBV   | cAg     | 141-151     | STLPETTVVRR<br>(SEQ. ID NO.:139)     | HLA-A*3101        | 132  |
| HBV   | cAg     | 141-151     | STLPETTVVRR<br>(SEQ. ID NO.:139)     | HLA-A*6801        | 132  |
| HBV   | cAg     | 18-27       | FLPSDFFPVS<br>(SEQ. ID NO.:126)      | HLA-A*6801        | 127  |
| HBV   | sAg     | 28-39       | IPQSLDSWWTSL<br>(SEQ. ID NO.:140)    | H2-L <sup>d</sup> | 133  |
| HBV   | cAg     | 93-100      | MGLKFRQL<br>(SEQ. ID NO.:141)        | H2-K <sup>b</sup> | 134  |
| HBV   | preS    | 141-149     | STBXQSGXQ<br>(SEQ. ID NO.:142)       | HLA-A*0201        | 135  |
| HCMV  | gp B    | 618-628     | FIAGNSAYEYV<br>(SEQ. ID NO.:143)     | HLA-A*0201        | 124  |
| HCMV  | E1      | 978-989     | SDEEEAIVAYTL<br>(SEQ. ID NO.:144)    | HLA-B18           | 136  |
| HCMV  | pp65    | 397-411     | DDVWTSGSDSDEELV<br>(SEQ. ID NO.:145) | HLA-b35           | 137  |
| HCMV  | pp65    | 123-131     | IPSINVHHY<br>(SEQ. ID NO.:146)       | HLA-B*3501        | 136  |
| HCMV  | pp65    | 495-504     | NLVPMVATVO<br>(SEQ. ID NO.:147)      | HLA-A*0201        | 137  |
| HCMV  | pp65    | 415-429     | RKTPTVTGGGAMAGA<br>(SEQ. ID NO.:148) | HLA-B7            | 137  |
| HCV   | MP      | 17-25       | DLMGYIPLV<br>(SEQ. ID NO.:149)       | HLA-A*0201        | 138  |
| HCV   | MP      | 63-72       | LLALLSCLTV<br>(SEQ. ID NO.:150)      | HLA-A*0201        | 139  |
| HCV   | MP      | 105-112     | ILHTPGCV<br>(SEQ. ID NO.:151)        | HLA-A*0201        | 139  |
| HCV   | env E   | 66-75       | QLRRHIDLLV<br>(SEQ. ID NO.:152)      | HLA-A*0201        | 139  |
| HCV   | env E   | 88-96       | DLCGSVFLV<br>(SEQ. ID NO.:153)       | HLA-A*0201        | 139  |
| HCV   | env E   | 172-180     | SMVGWAKV<br>(SEQ. ID NO.:154)        | HLA-A*0201        | 139  |
| HCV   | NS1     | 308-316     | HLHQNIVDV<br>(SEQ. ID NO.:155)       | HLA-A*0201        | 139  |
| HCV   | NS1     | 340-348     | FLLADARV<br>(SEQ. ID NO.:156)        | HLA-A*0201        | 139  |
| HCV   | NS2     | 234-246     | GLRDLAVAVEPVV<br>(SEQ. ID NO.:157)   | HLA-A*0201        | 139  |
| HCV   | NS1     | 18-28       | SLLAPGAKQNV<br>(SEQ. ID NO.:158)     | HLA-A*0201        | 139  |
| HCV   | NS1     | 19-28       | LLAPGAKQNV<br>(SEQ. ID NO.:159)      | HLA-A*0201        | 139  |

Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein      | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref. |
|-------|--------------|-------------|--------------------------------------|-------------------|------|
| HCV   | NS4          | 192-201     | LLFNILGGWV<br>(SEQ. ID NO.:160)      | HLA-A*0201        | 129  |
| HCV   | NS3          | 579-587     | YLVAYQATV<br>(SEQ. ID NO.:161)       | HLA-A*0201        | 129  |
| HCV   | core protein | 34-43       | YLLPRRGPRRL<br>(SEQ. ID NO.:162)     | HLA-A*0201        | 129  |
| HCV   | MP           | 63-72       | LLALLSCLTI<br>(SEQ. ID NO.:163)      | HLA-A*0201        | 129  |
| HCV   | NS4          | 174-182     | SLMAFTAADV<br>(SEQ. ID NO.:164)      | HLA-A*0201        | 140  |
| HCV   | NS3          | 67-75       | CINGVCWTV<br>(SEQ. ID NO.:165)       | HLA-A*0201        | 140  |
| HCV   | NS3          | 163-171     | LLCPAGHAV<br>(SEQ. ID NO.:166)       | HLA-A*0201        | 141  |
| HCV   | NS5          | 239-247     | ILDSFDPLV<br>(SEQ. ID NO.:167)       | HLA-A*0201        | 141  |
| HCV   | NS4A         | 236-244     | ILAGYGAGV<br>(SEQ. ID NO.:168)       | HLA-A*0201        | 142  |
| HCV   | NS5          | 714-722     | GLQDCTMLV<br>(SEQ. ID NO.:169)       | HLA-A*0201        | 142  |
| HCV   | NS3          | 281-290     | TGAPVTYSTY<br>(SEQ. ID NO.:170)      | HLA-A*0201        | 143  |
| HCV   | NS4A         | 149-157     | HMWNFISGI<br>(SEQ. ID NO.:171)       | HLA-A*0201        | 144  |
| HCV   | NS5          | 575-583     | RVCEKMALY<br>(SEQ. ID NO.:172)       | HLA-A*0201-A3     | 145  |
| HCV   | NS1          | 238-246     | TINYTIFK<br>(SEQ. ID NO.:173)        | HLA-A*1101        | 145  |
| HCV   | NS2          | 109-116     | YISWCLWW<br>(SEQ. ID NO.:174)        | HLA-A23           | 145  |
| HCV   | core protein | 40-48       | GPRLGVRAT<br>(SEQ. ID NO.:175)       | HLA-B7            | 145  |
| HIV-1 | gp120        | 380-388     | SFNCGGEFF<br>(SEQ. ID NO.:176)       | HLA-Cw*0401       | 185  |
| HIV-1 | RT           | 206-214     | TEMEKEGKI<br>(SEQ. ID NO.:177)       | H2-K <sup>k</sup> | 186  |
| HIV-1 | p17          | 18-26       | KIRLRPGGK<br>(SEQ. ID NO.:178)       | HLA-A*0301        | 187  |
| HIV-1 | p17          | 20-29       | RLRPGGKKKY<br>(SEQ. ID NO.:179)      | HLA-A*0301        | 188  |
| HIV-1 | RT           | 325-333     | AIFQSSMTK<br>(SEQ. ID NO.:180)       | HLA-A*0301        | 188  |
| HIV-1 | p17          | 84-92       | TLYCVHQRI<br>(SEQ. ID NO.:181)       | HLA-A11           | 188  |
| HIV-1 | RT           | 508-517     | IYQEPFKNLK<br>(SEQ. ID NO.:182)      | HLA-A11           | 188  |
| HIV-1 | p17          | 28-36       | KYKLKHIVW<br>(SEQ. ID NO.:183)       | HLA-A24           | 188  |

Table I: Viral epitopes on MHC class 1 molecules

| Virus | Protein | AA position | T cell epitope/MHC ligand (Antigen) | MHC molecule | Ref. |
|-------|---------|-------------|-------------------------------------|--------------|------|
| HIV-1 | gp120   | 53-62       | LFCASDAKAY<br>(SEQ. ID NO.:184)     | HLA-A24      | 189  |
| HIV-1 | gagp24  | 145-155     | QAISPRTLNAW<br>(SEQ. ID NO.:185)    | HLA-A25      | 188  |
| HIV-1 | gagp24  | 167-175     | EVIPMFSAAL<br>(SEQ. ID NO.:186)     | HLA-A26      | 188  |
| HIV-1 | RT      | 593-603     | ETFYVDGAANR<br>(SEQ. ID NO.:187)    | HLA-A26      | 188  |
| HIV-1 | gp41    | 775-785     | RLRDLLIVTR<br>(SEQ. ID NO.:188)     | HLA-A31      | 190  |
| HIV-1 | RT      | 559-568     | PIQKETWETW<br>(SEQ. ID NO.:189)     | HLA-A32      | 187  |
| HIV-1 | gp120   | 419-427     | RIKQINMW<br>(SEQ. ID NO.:190)       | HLA-A32      | 187  |
| HIV-1 | RT      | 71-79       | ITLWQRPLV<br>(SEQ. ID NO.:191)      | HLA-A*6802   | 188  |
| HIV-1 | RT      | 85-93       | DTVLEEMNL<br>(SEQ. ID NO.:192)      | HLA-A*6802   | 188  |
| HIV-1 | RT      | 71-79       | ITLWQRPLV<br>(SEQ. ID NO.:193)      | HLA-A*7401   | 188  |
| HIV-1 | gag p24 | 148-156     | SPRTLNAWV<br>(SEQ. ID NO.:194)      | HLA-B7       | 188  |
| HIV-1 | gagp24  | 179-187     | ATPQDLNTM<br>(SEQ. ID NO.:195)      | HLA-B7       | 188  |
| HIV-1 | gp120   | 303-312     | RPNNNTRKSI<br>(SEQ. ID NO.:196)     | HLA-B7       | 188  |
| HIV-1 | gp41    | 843-851     | IPRRIROGL<br>(SEQ. ID NO.:197)      | HLA-B7       | 188  |
| HIV-1 | p17     | 74-82       | ELRSLYNTV<br>(SEQ. ID NO.:198)      | HLA-B8       | 188  |
| HIV-1 | nef     | 13-20       | WPTVRERM<br>(SEQ. ID NO.:199)       | HLA-B8       | 188  |
| HIV-1 | nef     | 90-97       | FLKEKGGL<br>(SEQ. ID NO.:200)       | HLA-B8       | 188  |
| HIV-1 | gag p24 | 183-191     | DLNTMLNTV<br>(SEQ. ID NO.:568)      | HLA-B14      | 191  |
| HIV-1 | P17     | 18-27       | KIRLRPGGKK<br>(SEQ. ID NO.:201)     | HLA-B27      | 188  |
| HIV-1 | p17     | 19-27       | IRLRPGGKK<br>(SEQ. ID NO.:202)      | HLA-B27      | 188  |
| HIV-1 | gp41    | 791-799     | GRRGWEALKY<br>(SEQ. ID NO.:203)     | HLA-B27      | 188  |
| HIV-1 | nef     | 73-82       | QVPLRPMTYK<br>(SEQ. ID NO.:204)     | HLA-B27      | 188  |
| HIV-1 | GP41    | 590-597     | RYLKDQQL<br>(SEQ. ID NO.:205)       | HLA-B27      | 192  |
| HIV-1 | nef     | 105-114     | RRQDILDWL<br>(SEQ. ID NO.:206)      | HLA-B*2705   | 188  |

**Table I: Viral epitopes on MHC class 1 molecules**

| <b>Virus</b> | <b>Protein</b> | <b>AA position</b> | <b>T cell epitope/ MHC ligand (Antigen)</b> | <b>MHC molecule</b> | <b>Ref.</b> |
|--------------|----------------|--------------------|---|---------------------|-------------|
| HIV-1        | nef            | 134-141            | RYPLTFGW<br>(SEQ. ID NO.:207)               | HLA-B*2705          | 188         |
| HIV-1        | p17            | 36-44              | WASRELERF<br>(SEQ. ID NO.:208)              | HLA-B35             | 188         |
| HIV-1        | GAG P24        | 262-270            | TVLDVGDY<br>(SEQ. ID NO.:209)               | HLA-B35             | 188         |
| HIV-1        | gp120          | 42-52              | VPVWKEATTTL<br>(SEQ. ID NO.:210)            | HLA-B35             | 188         |
| HIV-1        | P17            | 36-44              | NSSKVSQNY<br>(SEQ. ID NO.:221)              | HLA-B35             | 193         |
| HIV-1        | gag p24        | 254-262            | PPIPVGDIY<br>(SEQ. ID NO.:212)              | HLA-B35             | 193         |
| HIV-1        | RT             | 342-350            | HPDIVIYQY<br>(SEQ. ID NO.:213)              | HLA-B35             | 193         |
| HIV-1        | gp41           | 611-619            | TAVPWNASW<br>(SEQ. ID NO.:214)              | HLA-B35             | 194         |
| HIV-1        | gag            | 245-253            | NPVPVGNIIY<br>(SEQ. ID NO.:215)             | HLA-B35             | 193         |
| HIV-1        | nef            | 120-128            | YFPDWQNYT<br>(SEQ. ID NO.:216)              | HLA-B37             | 188         |
| HIV-1        | gag p24        | 193-201            | GHQAAMQML<br>(SEQ. ID NO.:217)              | HLA-B42             | 188         |
| HIV-1        | p17            | 20-29              | RLRPGGKKKY<br>(SEQ. ID NO.:218)             | HLA-B42             | 188         |
| HIV-1        | RT             | 438-446            | YPGIKVRQL<br>(SEQ. ID NO.:219)              | HLA-B42             | 188         |
| HIV-1        | RT             | 591-600            | GAETFYVDGA<br>(SEQ. ID NO.:220)             | HLA-B45             | 188         |
| HIV-1        | gag p24        | 325-333            | NANPDCKTI<br>(SEQ. ID NO.:221)              | HLA-B51             | 188         |
| HIV-1        | gag p24        | 275-282            | RMYSPTSI<br>(SEQ. ID NO.:222)               | HLA-B52             | 188         |
| HIV-1        | gp120          | 42-51              | VPVWKEATTT<br>(SEQ. ID NO.:223)             | HLA-B*5501          | 192         |
| HIV-1        | gag p24        | 147-155            | ISPRTLNAW<br>(SEQ. ID NO.:224)              | HLA-B57             | 188         |
| HIV-1        | gag p24        | 240-249            | TSTLQEQIGW<br>(SEQ. ID NO.:225)             | HLA-B57             | 188         |
| HIV-1        | gag p24        | 162-172            | KAFSPEVIPMF<br>(SEQ. ID NO.:226)            | HLA-B57             | 188         |
| HIV-1        | gag p24        | 311-319            | QASQEVKNW<br>(SEQ. ID NO.:227)              | HLA-B57             | 188         |
| HIV-1        | gag p24        | 311-319            | QASQDVKNW<br>(SEQ. ID NO.:228)              | HLA-B57             | 188         |
| HIV-1        | nef            | 116-125            | HTQGYFPDWQ<br>(SEQ. ID NO.:229)             | HLA-B57             | 188         |
| HIV-1        | nef            | 120-128            | YFPDWQNYT<br>(SEQ. ID NO.:230)              | HLA-B57             | 188         |

Table I: Viral epitopes on MHC class 1 molecules

| Virus                | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref. |
|----------------------|---------|-------------|--------------------------------------|-------------------|------|
| HIV-1                | gag p24 | 240-249     | TSTLQEQIGW<br>(SEQ. ID NO.:231)      | HLA-B58           | 188  |
| HIV-1                | p17     | 20-29       | RLRPGGKKKY<br>(SEQ. ID NO.:232)      | HLA-B62           | 188  |
| HIV-1                | p24     | 268-277     | LGLNKIVRMY<br>(SEQ. ID NO.:233)      | HLA-B62           | 188  |
| HIV-1                | RT      | 415-426     | LVGKLNWASQIY<br>(SEQ. ID NO.:234)    | HLA-B62           | 188  |
| HIV-1                | RT      | 476-485     | ILKEPVHGVY<br>(SEQ. ID NO.:235)      | HLA-B62           | 188  |
| HIV-1                | nef     | 117-127     | TQGYFPDWQNY<br>(SEQ. ID NO.:236)     | HLA-B62           | 188  |
| HIV-1                | nef     | 84-91       | AVDLSHFL<br>(SEQ. ID NO.:237)        | HLA-B62           | 188  |
| HIV-1                | gag p24 | 168-175     | VIPMFSA<br>(SEQ. ID NO.:238)         | HLA-Cw*0102       | 188  |
| HIV-1                | gp120   | 376-384     | FNCGGEFFY<br>(SEQ. ID NO.:239)       | HLA-A29           | 196  |
| HIV-1                | gp 120  | 375-383     | SFNCGGEFF<br>(SEQ. ID NO.:240)       | HLA-B15           | 196  |
| HIV-1                | nef     | 136-145     | PLTFGWCYKL<br>(SEQ. ID NO.:241)      | HLA-A*0201        | 197  |
| HIV-1                | nef     | 180-189     | VLEWRFSRL<br>(SEQ. ID NO.:242)       | HLA-A*0201        | 197  |
| HIV-1                | nef     | 68-77       | FPVTPQVPLR<br>(SEQ. ID NO.:243)      | HLA-B7            | 197  |
| HIV-1                | nef     | 128-137     | TPGPGVRYPL<br>(SEQ. ID NO.:244)      | HLA-B7            | 197  |
| HIV-1                | gag p24 | 308-316     | QASQEVKNW<br>(SEQ. ID NO.:245)       | HLA-Cw*0401       | 521  |
| HIV-1 IIIB           | RT      | 273-282     | VPLDEDFRKY<br>(SEQ. ID NO.:246)      | HLA-B35           | 181  |
| HIV-1 IIIB           | RT      | 25-33       | NPDIVIYQY<br>(SEQ. ID NO.:247)       | HLA-B35           | 181  |
| HIV-1 IIIB           | gp41    | 557-565     | RAIEAQAHL<br>(SEQ. ID NO.:248)       | HLA-B51           | 181  |
| HIV-1 IIIB           | RT      | 231-238     | TAFTIPSI<br>(SEQ. ID NO.:249)        | HLA-B51           | 181  |
| HIV-1 IIIB           | p24     | 215-223     | VHPVHAGPIA<br>(SEQ. ID NO.:250)      | HLA-B*5501        | 181  |
| HIV-1 IIIB           | gp120   | 156-165     | NCSFNISTSI<br>(SEQ. ID NO.:251)      | HLA-Cw8           | 181  |
| HIV-1 IIIB           | gp120   | 241-249     | CTNVSTVQC<br>(SEQ. ID NO.:252)       | HLA-Cw8           | 181  |
| HIV-1 <sub>5F2</sub> | gp120   | 312-320     | IGPGRAFHT<br>(SEQ. ID NO.:253)       | H2-D <sup>d</sup> | 198  |
| HIV-1 <sub>5F2</sub> | pol     | 25-33       | NPDIVIYQY<br>(SEQ. ID NO.:254)       | HLA-B*3501        | 199  |

Table I: Viral epitopes on MHC class 1 molecules

| Virus                | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref. |
|----------------------|---------|-------------|--------------------------------------|-------------------|------|
| HIV-1 <sub>5F2</sub> | pol     | 432-441     | EPIVGAETFY<br>(SEQ. ID NO.:255)      | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | pol     | 432-440     | EPIVGAETF<br>(SEQ. ID NO.:256)       | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | pol     | 6-14        | SPAIFQSSM<br>(SEQ. ID NO.:257)       | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | pol     | 59-68       | VPLDKDFRKY<br>(SEQ. ID NO.:258)      | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | pol     | 6-14        | IPLTEEAEL<br>(SEQ. ID NO.:259)       | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | nef     | 69-79       | RPQVPLRPMTY<br>(SEQ. ID NO.:260)     | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | nef     | 66-74       | FPVRPQVPL<br>(SEQ. ID NO.:261)       | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | env     | 10-18       | DPNPQEVVL<br>(SEQ. ID NO.:262)       | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | env     | 7-15        | RPIVSTQLL<br>(SEQ. ID NO.:263)       | HLA-B*3501        | 199  |
| HIV-1 <sub>5F2</sub> | pol     | 6-14        | IPLTEEAEL<br>(SEQ. ID NO.:264)       | HLA-B51           | 199  |
| HIV-1 <sub>5F2</sub> | env     | 10-18       | DPNPQEVVL<br>(SEQ. ID NO.:265)       | HLA-B51           | 199  |
| HIV-1 <sub>5F2</sub> | gagp24  | 199-207     | AMQMLKETI<br>(SEQ. ID NO.:266)       | H2-K <sup>d</sup> | 198  |
| HIV-2                | gagp24  | 182-190     | TPYDINQML<br>(SEQ. ID NO.:267)       | HLA-B*5301        | 200  |
| HIV-2                | gag     | 260-269     | RRWQLGLQKV<br>(SEQ. ID NO.:268)      | HLA-B*2703        | 188  |
| HIV-1 <sub>5F2</sub> | gp41    | 593-607     | GIWGCSGKLICTAV<br>(SEQ. ID NO.:269)  | HLA-B17           | 201  |
| HIV-1 <sub>5F2</sub> | gp41    | 753-767     | ALIWEDLRSLCLFSY<br>(SEQ. ID NO.:270) | HLA-B22           | 201  |
| HPV 6b               | E7      | 21-30       | GLHCYEQLV<br>(SEQ. ID NO.:271)       | HLA-A*0201        | 202  |
| HPV 6b               | E7      | 47-55       | PLKQHFQIV<br>(SEQ. ID NO.:272)       | HLA-A*0201        | 202  |
| HPV11                | E7      | 4-12        | RLVTLKDIV<br>(SEQ. ID NO.:273)       | HLA-A*0201        | 202  |
| HPV16                | E7      | 86-94       | TLGIVCPIC<br>(SEQ. ID NO.:274)       | HLA-A*0201        | 129  |
| HPV16                | E7      | 85-93       | GTLGIVCPI<br>(SEQ. ID NO.:275)       | HLA-A*0201        | 129  |
| HPV16                | E7      | 12-20       | MLDLQPETT<br>(SEQ. ID NO.:276)       | HLA-A*0201        | 129  |
| HPV16                | E7      | 11-20       | YMLDLQPETT<br>(SEQ. ID NO.:277)      | HLA-A*0201        | 203  |
| HPV16                | E6      | 15-22       | RPRKLPQL<br>(SEQ. ID NO.:278)        | HLA-B7            | 204  |



Table I: Viral epitopes on MHC class 1 molecules

| Virus       | Protein | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref.                      |
|-------------|---------|-------------|--------------------------------------|-------------------|---------------------------|
| HPV16e6     | 15-22   | 49-57       | RAHYNIVTF<br>(SEQ. ID NO.:279)       | HW-D <sup>b</sup> | 205                       |
| HSV         | gp B    | 498-505     | SSIEFARL<br>(SEQ. ID NO.:280)        | H2-K <sup>b</sup> | 206                       |
| HSV-1       | gp C    | 480-488     | GIGIGVLAA<br>(SEQ. ID NO.:281)       | HLA-A*0201        | 104                       |
| HSV-1       | ICP27   | 448-456     | DYATLGVG<br>(SEQ. ID NO.:282)        | H2-K <sup>d</sup> | 207                       |
| HSV-1       | ICP27   | 322-332     | LYRTFAGNPRA<br>(SEQ. ID NO.:283)     | H2-K <sup>d</sup> | 207                       |
| HSV-1       | UL39    | 822-829     | QTFDFGRL<br>(SEQ. ID NO.:284)        | H2-K <sup>b</sup> | 208                       |
| HSV-2       | gp C    | 446-454     | GAGIGVAVL<br>(SEQ. ID NO.:285)       | HLA-A*0201        | 104                       |
| HTLV-1      | TAX     | 11-19       | LLFGYPVYV<br>(SEQ. ID NO.:286)       | HLA-A*0201        | 209                       |
| Influenza   | MP      | 58-66       | GILGFVFTL<br>(SEQ. ID NO.:287)       | HLA-A*0201        | 68, 169, 209,<br>210, 211 |
| Influenza   | MP      | 59-68       | ILGFVFTLTV<br>(SEQ. ID NO.:288)      | HLA-A*0201        | 168, 212, 213             |
| Influenza   | NP      | 265-273     | ILRGSAVHK<br>(SEQ. ID NO.:289)       | HLA-A3            | 214                       |
| Influenza   | NP      | 91-99       | KTGGPIYKR<br>(SEQ. ID NO.:290)       | HLA-A*6801        | 215, 216                  |
| Influenza   | NP      | 380-388     | ELRSRYWAI<br>(SEQ. ID NO.:291)       | HLA-B8            | 217                       |
| Influenza   | NP      | 381-388     | LRSRYWAI<br>(SEQ. ID NO.:292)        | HLA-B*2702        | 218                       |
| Influenza   | NP      | 339-347     | EDLRVLSFI<br>(SEQ. ID NO.:293)       | HLA-B*3701        | 219                       |
| Influenza   | NS1     | 158-166     | GEISPLPSL<br>(SEQ. ID NO.:294)       | HLA-B44           | 220                       |
| Influenza   | NP      | 338-346     | FEDLRVLSF<br>(SEQ. ID NO.:295)       | HLA-B44           | 220                       |
| Influenza   | NS1     | 158-166     | GEISPLPSL<br>(SEQ. ID NO.:294)       | HLA-B*4402        | 220                       |
| Influenza   | NP      | 338-346     | FEDLRVLSF<br>(SEQ. ID NO.:295)       | HLA-B*4402        | 220                       |
| Influenza   | PB1     | 591-599     | VSDGGPNLY<br>(SEQ. ID NO.:296)       | HLA-A1            | 214, 29                   |
| Influenza A | NP      | 44-52       | CTELKLSDY<br>(SEQ. ID NO.:297)       | HLA-A1            | 29                        |
| Influenza   | NS1     | 122-130     | AIMDKNIL<br>(SEQ. ID NO.:298)        | HLA-A*0201        | 221                       |
| Influenza A | NS1     | 123-132     | IMDKNILKA<br>(SEQ. ID NO.:299)       | HLA-A*0201        | 221                       |
| Influenza A | NP      | 383-391     | SRYWAIATR<br>(SEQ. ID NO.:300)       | HLA-B*2705        | 160, 184                  |

Table I: Viral epitopes on MHC class 1 molecules

| Virus         | Protein       | AA position | T cell epitope/MHC ligand (Antigen)   | MHC molecule      | Ref           |
|---------------|---------------|-------------|---------------------------------------|-------------------|---------------|
| Influenza A   | NP            | 147-155     | TYQRTALV<br>(SEQ. ID NO.:301)         | H2-K <sup>d</sup> | 222, 223      |
| Influenza A   | HA            | 210-219     | TYVSVSTSL<br>(SEQ. ID NO.:302)        | H2-K <sup>d</sup> | 224, 225      |
| Influenza A   | HA            | 518-526     | IYSTVASSL<br>(SEQ. ID NO.:303)        | H2-K <sup>d</sup> | 224           |
| Influenza A   | HA            | 259-266     | FEANGNLI<br>(SEQ. ID NO.:304)         | H2-K <sup>k</sup> | 226, 227, 228 |
| Influenza A   | HA            | 10-18       | IEGGWTGMI<br>(SEQ. ID NO.:305)        | H2-K <sup>k</sup> | 226, 227, 228 |
| Influenza A   | NP            | 50-57       | SDYEGRLI<br>(SEQ. ID NO.:306)         | H2-K <sup>k</sup> | 229, 230      |
| Influenza a   | NS1           | 152-160     | EEGAIVGEI<br>(SEQ. ID NO.:307)        | H2-K <sup>k</sup> | 231           |
| Influenza A34 | NP            | 366-374     | ASNENMETM<br>(SEQ. ID NO.:308)        | H2-D <sup>b</sup> | 168, 222, 219 |
| Influenza A68 | NP            | 366-374     | ASNENMDAM<br>(SEQ. ID NO.:309)        | H2-D <sup>b</sup> | 232           |
| Influenza B   | NP            | 85-94       | KLGEFYNQMM<br>(SEQ. ID NO.:310)       | HLA-A*0201        | 233           |
| Influenza B   | NP            | 85-94       | KAGEFYNQMM<br>(SEQ. ID NO.:311)       | HLA-A*0201        | 234           |
| Influenza JAP | HA            | 204-212     | LYQNVGTYV<br>(SEQ. ID NO.:312)        | H2-K <sup>d</sup> | 235           |
| Influenza JAP | HA            | 210-219     | TYVSVGTSL<br>(SEQ. ID NO.:313)        | H2-K <sup>d</sup> | 225           |
| Influenza JAP | HA            | 523-531     | VYQILAIYA<br>(SEQ. ID NO.:314)        | H2-K <sup>d</sup> | 235           |
| Influenza JAP | HA            | 529-537     | IYATVAGSL<br>(SEQ. ID NO.:315)        | H2-K <sup>d</sup> | 235           |
| Influenza JAP | HA            | 210-219     | TYVSVGTSTI (L>I)<br>(SEQ. ID NO.:316) | H2-K <sup>d</sup> | 236           |
| Influenza JAP | HA            | 255-262     | FESTGNLI<br>(SEQ. ID NO.:317)         | H2-K <sup>k</sup> | 237           |
| JHMOV         | cAg           | 318-326     | APTAGAFFF<br>(SEQ. ID NO.:318)        | H2-L <sup>d</sup> | 238           |
| LCMV          | NP            | 118-126     | RPQASGVYM<br>(SEQ. ID NO.:319)        | H2-L <sup>d</sup> | 239, 240      |
| LCMV          | NP            | 396-404     | FQPQNGQFI<br>(SEQ. ID NO.:320)        | H2-D <sup>b</sup> | 241           |
| LCMV          | GP            | 276-286     | SGVENPGGYCL<br>(SEQ. ID NO.:321)      | H2-D <sup>b</sup> | 242           |
| LCMV          | GP            | 33-42       | KAVYNFATCG<br>(SEQ. ID NO.:322)       | H2-D <sup>b</sup> | 243, 244      |
| MCMV          | pp89          | 168-176     | YPHFMPITNL<br>(SEQ. ID NO.:323)       | H2-L <sup>d</sup> | 245           |
| MHV           | spike protein | 510-518     | CLSWNGPHL<br>(SEQ. ID NO.:324)        | H2-D <sup>b</sup> | 248           |

Table I: Viral epitopes on MHC class 1 molecules

| Virus                    | Protein   | AA position | T cell epitope/ MHC ligand (Antigen) | MHC molecule      | Ref.                           |
|--------------------------|-----------|-------------|--------------------------------------|-------------------|--------------------------------|
| MMTV                     | env gp 36 | 474-482     | SFAVATTAL<br>(SEQ. ID NO.:325)       | H2-K <sup>d</sup> | 246                            |
| MMTV                     | gag p27   | 425-433     | SYETFISRL<br>(SEQ. ID NO.:326)       | H2-K <sup>d</sup> | 246                            |
| MMTV                     | env gp73  | 544-551     | ANYDFICV<br>(SEQ. ID NO.:327)        | H2-K <sup>b</sup> | 247                            |
| MuLV                     | env p15E  | 574-581     | KSPWFTTL<br>(SEQ. ID NO.:328)        | H2-K <sup>b</sup> | 249, 250                       |
| MuLV                     | env gp70  | 189-196     | SSWDFITV<br>(SEQ. ID NO.:329)        | H2-K <sup>b</sup> | 251, Sijts et al.<br>submitted |
| MuLV                     | gag 75k   | 75-83       | CCLCLTVFL<br>(SEQ. ID NO.:330)       | H2-D <sup>b</sup> | 252                            |
| MuLV                     | env gp70  | 423-431     | SPSYVYHQF<br>(SEQ. ID NO.:331)       | H2-L <sup>d</sup> | 253                            |
| MV                       | F protein | 437-447     | SRRYPDAVYLH<br>(SEQ. ID NO.:332)     | HLA-B*27          | 254                            |
| MV                       | F protein | 438-446     | RRYPDVAVYL<br>(SEQ. ID NO.:333)      | HLA-B*2705        | 255                            |
| MV                       | NP        | 281-289     | YPALGLHEF<br>(SEQ. ID NO.:334)       | H2-L <sup>d</sup> | 256                            |
| MV                       | HA        | 343-351     | DPVIDRLYL<br>(SEQ. ID NO.:335)       | H2-L <sup>d</sup> | 257                            |
| MV                       | HA        | 544-552     | SPGRSFSYF<br>(SEQ. ID NO.:336)       | H2-L <sup>d</sup> | 257                            |
| Poliovirus               | VP1       | 111-118     | TYKDTVQL<br>(SEQ. ID NO.:337)        | H2-K <sup>d</sup> | 258                            |
| Poliovirus               | VP1       | 208-217     | FYDGFSGVPL<br>(SEQ. ID NO.:338)      | H2-K <sup>d</sup> | 258                            |
| Pseudorabies<br>virus gp | G111      | 455-463     | IAGIGILAI<br>(SEQ. ID NO.:339)       | HLA-A*0201        | 104                            |
| Rabiesvirus              | NS        | 197-205     | VEAEIAHQI<br>(SEQ. ID NO.:340)       | H2-K <sup>k</sup> | 227, 227                       |
| Rotavirus                | VP7       | 33-40       | IIYRFLLI<br>(SEQ. ID NO.:341)        | H2-K <sup>b</sup> | 259                            |
| Rotavirus                | VP6       | 376-384     | VGPVFPFGM<br>(SEQ. ID NO.:342)       | H2-K <sup>b</sup> | 260                            |
| Rotavirus                | VP3       | 585-593     | YSGYIFRDL<br>(SEQ. ID NO.:343)       | H2-K <sup>b</sup> | 260                            |
| RSV                      | M2        | 82-90       | SYIGSINNI<br>(SEQ. ID NO.:344)       | H2-K <sup>d</sup> | 261                            |
| SIV                      | gagp11C   | 179-190     | EGCTPYDINQML<br>(SEQ. ID NO.:345)    | Mamu-A*01         | 266                            |
| SV                       | NP        | 324-332     | FAPGNYPAL<br>(SEQ. ID NO.:346)       | H2-D <sup>b</sup> | 262                            |
| SV                       | NP        | 324-332     | FAPGNYPAL<br>(SEQ. ID NO.:346)       | H2-K <sup>b</sup> | 263, 264, 265                  |

**Table I: Viral epitopes on MHC class 1 molecules**

| <b>Virus</b> | <b>Protein</b> | <b>AA position</b> | <b>T cell epitope/ MHC ligand (Antigen)</b> | <b>MHC molecule</b> | <b>Ref.</b> |
|--------------|----------------|--------------------|---|---------------------|-------------|
| SV40         | T              | 404-411            | VVYDFLKC<br>(SEQ. ID NO.:347)               | H2-K <sup>b</sup>   | 267         |
| SV40         | T              | 206-215            | SAINNYAQKL<br>(SEQ. ID NO.:348)             | H2-D <sup>b</sup>   | 268, 269    |
| SV40         | T              | 223-231            | CKGVNKEYL<br>(SEQ. ID NO.:349)              | H2-D <sup>b</sup>   | 268, 269    |
| SV40         | T              | 489-497            | QGINNLDNL<br>(SEQ. ID NO.:350)              | H2-D <sup>b</sup>   | 268, 269    |
| SV40         | T              | 492-500<br>(501)   | NNLDNLRDY(L)<br>(SEQ. ID NO.:351)           | H2-D <sup>b</sup>   | 270         |
| SV40         | T              | 560-568            | SEFLLEKRI<br>(SEQ. ID NO.:352)              | H2-K <sup>k</sup>   | 271         |
| VSV          | NP             | 52-59              | RGYVYQGL<br>(SEQ. ID NO.:353)               | H2-K <sup>b</sup>   | 272         |

Table II sets forth antigens identified from various protein sources. The Table is extracted from Table 4.2 in the Rammansee book with the references in Table II being the same as the references in the Rammensee Table 4.2.

| TABLE II<br>HLA Class I Motifs |  |                                   |             |
|--------------------------------|--|-----------------------------------|-------------|
| HLA -A1                        | Position (Antigen)                         | Source                            | Ref.        |
| T cell epitopes                | E A D P T G H S Y<br>(SEQ. ID NO.:354)     | MAGE-1 161-169                    | 27,28       |
|                                | V S D G G P N L Y<br>(SEQ. ID NO.:355)     | Influenza A PB1 591-599           | 21,23       |
|                                | C T E L K L S D Y<br>(SEQ. ID NO.:356)     | Influenza A NP 44-52              | 23          |
|                                | E V D P I G H L Y<br>(SEQ. ID NO.:357)     | MAGE-3 168-176                    | 29,30       |
| HLA-A201                       | M L L S V P L L L G<br>(SEQ. ID NO.:358)   | Calreticulin signal sequence 1-10 | 34,35,36,37 |
|                                | S T B X Q S G X Q<br>(SEQ. ID NO.:359)     | HBV PRE-S PROTEIN 141-149         | 43          |
|                                | Y M D G T M S Q V<br>(SEQ. ID NO.:360)     | Tyrosinase 369-377                | 45          |
|                                | I L K E P V H G V<br>(SEQ. ID NO.:361)     | HIV-1 RT 476-484                  | 4,31,47     |
|                                | I L G F V F T L T V<br>(SEQ. ID NO.:362)   | Influenza MP 59-68                | 4,39        |
|                                | L L F G Y P V Y V V<br>(SEQ. ID NO.:363)   | HTLV -I tax 11-19                 | 40          |
|                                | G L S P T V W L S V<br>(SEQ. ID NO.:364)   | HBV sAg 348-357                   | 48          |
|                                | W L S L L V P F V<br>(SEQ. ID NO.:365)     | HBV sAg 335-343                   | 49,50,51    |
|                                | F L P S D F F P S V<br>(SEQ. ID NO.:366)   | HBV cAg 18-27                     | 52          |
|                                | C L G G L L T M V<br>(SEQ. ID NO.:367)     | EBV LMP-2 426-434                 | 48          |
|                                | F L A G N S A Y E Y V<br>(SEQ. ID NO.:368) | HCMV gp 618-628B                  | 53          |

**TABLE II**  
**HLA Class I Motifs**

|  |                           |             |
|--|---------------------------|-------------|
| K L G E F Y N Q M M<br>(SEQ. ID NO.:369) | Influenza BNP 85-94       | 54          |
| K L V A L G I N A V<br>(SEQ. ID NO.:370) | HCV-1 NS3 400-409         | 55          |
| D L M G Y I P L V<br>(SEQ. ID NO.:371)   | HCV MP 17-25              | 56          |
| R L V T L K D I V<br>(SEQ. ID NO.:372)   | HPV11 EZ 4-12             | 34,35       |
| M L L A V L Y C L<br>(SEQ. ID NO.:373)   | Tyrosinase 1-9            | 57,58,59,68 |
| A A G I G I L T V<br>(SEQ. ID NO.:374)   | Melan A/Mart-127-35       | 60          |
| Y L L P G P V T A<br>(SEQ. ID NO.:375)   | Pmel 17/gp100 480-488     | 61          |
| I I D G T A T L R L<br>(SEQ. ID NO.:376) | Pmel 17/ gp100 457-466    | 62          |
| I L D G T A T L R L<br>(SEQ. ID NO.:377) | Pmel gp100 457-466        | 62          |
| I T D Q V P F S V<br>(SEQ. ID NO.:378)   | Pmel gp100 209-217        | 62          |
| K I W G Q Y W Q V<br>(SEQ. ID NO.:379)   | Pmel gp100 154-162        | 62          |
| T I T D Q V P F S V<br>(SEQ. ID NO.:380) | Pmel gp100 208-217        | 62          |
| A F H H V A R E L<br>(SEQ. ID NO.:381)   | HIV-1 nef 190-198         | 63          |
| Y L N K I Q N S L<br>(SEQ. ID NO.:382)   | P. falciparum CSP 334-342 | 64          |
| M M R K L A I L S V<br>(SEQ. ID NO.:383) | P. falciparum CSP 1-10    | 64          |
| K A G E F Y N Q M M<br>(SEQ. ID NO.:384) | Influenza BNP 85-94       | 65          |
| N I A E G L R A L<br>(SEQ. ID NO.:385)   | EBNA-1 480-488            | 66          |
| N L R R G T A L A<br>(SEQ. ID NO.:386)   | EBNA-1 519-527            | 66          |

TABLE II  
HLA Class I Motifs

|                                 |                |    |
|---------------------------------|----------------|----|
| ALAIPQCRL<br>(SEQ. ID NO.:387)  | EBNA-1 525-533 | 66 |
| VLKDAIKDL<br>(SEQ. ID NO.:388)  | EBNA-1 575-582 | 66 |
| FMVFLQTHI<br>(SEQ. ID NO.:389)  | EBNA1 562-570  | 66 |
| HLIVDTDSL<br>(SEQ. ID NO.:390)  | EBNA-2 15-23   | 66 |
| SLGNPSLSV<br>(SEQ. ID NO.:391)  | EBNA-2 22-30   | 66 |
| PLASAMRML<br>(SEQ. ID NO.:392)  | EBNA-2 126-134 | 66 |
| RMLWMANYI<br>(SEQ. ID NO.:393)  | EBNA-2 132-140 | 66 |
| MLWMANYIV<br>(SEQ. ID NO.:394)  | EBNA-2 133-141 | 66 |
| ILPQGPQTA<br>(SEQ. ID NO.:395)  | EBNA-2 151-159 | 66 |
| PLRPTAPTTI<br>(SEQ. ID NO.:396) | EBNA-2 171-179 | 66 |
| PLPPATLTV<br>(SEQ. ID NO.:397)  | EBNA-2 205-213 | 66 |
| RMHLPVLHV<br>(SEQ. ID NO.:397)  | EBNA-2 246-254 | 66 |
| PMPLPPSQL<br>(SEQ. ID NO.:399)  | EBNA-2 287-295 | 66 |
| QLPPPAAPA<br>(SEQ. ID NO.:400)  | EBNA-2 294-302 | 66 |
| SMPELSPVL<br>(SEQ. ID NO.:401)  | EBNA-2 381-389 | 66 |
| DLDESWDYI<br>(SEQ. ID NO.:402)  | EBNA-2 453-461 | 66 |
| PLPCVLWPVV<br>(SEQ. ID NO.:403) | BZLF1 43-51    | 66 |
| SLEECDSEL<br>(SEQ. ID NO.:404)  | BZLF1 167-175  | 66 |

TABLE II  
HLA Class I Motifs

|                                    |                         |    |
|------------------------------------|-------------------------|----|
| EIKRYKNRV<br>(SEQ. ID NO.:405)     | BZLF1 176-184           | 66 |
| QLLQHYREV<br>(SEQ. ID NO.:406)     | BZLF1 195-203           | 66 |
| LLQHYREVA<br>(SEQ. ID NO.:407)     | BZLF1 196-204           | 66 |
| LLKQMCPSL<br>(SEQ. ID NO.:408)     | BZLF1 217-225           | 66 |
| SIIPRTPDV<br>(SEQ. ID NO.:409)     | BZLF1 229-237           | 66 |
| AIMDKNIIL<br>(SEQ. ID NO.:410)     | Influenza A NS1 122-130 | 67 |
| IMDKNIILKA<br>(SEQ. ID NO.:411)    | Influenza A NS1 123-132 | 67 |
| LLALLSCLTV<br>(SEQ. ID NO.:412)    | HCV MP 63-72            | 69 |
| ILHTPGCV<br>(SEQ. ID NO.:413)      | HCV MP 105-112          | 69 |
| QLRRHIDLLV<br>(SEQ. ID NO.:414)    | HCV env E 66-75         | 69 |
| DLCGSVFLV<br>(SEQ. ID NO.:415)     | HCV env E 88-96         | 69 |
| SMVG NWAKV<br>(SEQ. ID NO.:416)    | HCV env E 172-180       | 69 |
| HLHQNIVDV<br>(SEQ. ID NO.:417)     | HCV NS1 308-316         | 69 |
| FLLLADARV<br>(SEQ. ID NO.:418)     | HCV NS1 340-348         | 69 |
| GLRDLAVAVEPVV<br>(SEQ. ID NO.:419) | HCV NS2 234-246         | 69 |
| SLLAPGAKQNV<br>(SEQ. ID NO.:420)   | HCV NS1 18-28           | 69 |
| LLAPGAKQNV<br>(SEQ. ID NO.:421)    | HCV NS1 19-28           | 69 |
| FLLSLGIHL<br>(SEQ. ID NO.:422)     | HBV pol 575-583         | 70 |



**TABLE II**  
**HLA Class I Motifs**

|                                 |  |    |
|---------------------------------|--|----|
| SLYADSPSV<br>(SEQ. ID NO.:423)  | HBV pol 816-824  | 70 |
| GLSRYVARL<br>(SEQ. ID NO.:424)  | HBV POL 455-463  | 70 |
| KIFGSLAFL<br>(SEQ. ID NO.:425)  | HER-2 369-377  | 71 |
| ELVSEFSRM<br>(SEQ. ID NO.:426)  | HER-2 971-979  | 71 |
| KLTPLCVTL<br>(SEQ. ID NO.:427)  | HIV-1 gp160 120-128                                    | 72 |
| SLLNATDIAV<br>(SEQ. ID NO.:428) | HIV-1 GP160 814-823                                    | 72 |
| VLYRYGSFSV<br>(SEQ. ID NO.:429) | Pmel gp100 476-485                                     | 62 |
| YIGEVLVSV<br>(SEQ. ID NO.:430)  | Non-filament forming class I myosin<br>family (HA-2)** | 73 |
| LLFNILGGWV<br>(SEQ. ID NO.:431) | HCV NS4 192-201  | 74 |
| LLVPFVQWFW<br>(SEQ. ID NO.:432) | HBV env 338-347  | 74 |
| ALMPLYACI<br>(SEQ. ID NO.:433)  | HBV pol 642-650  | 74 |
| YLVAYQATV<br>(SEQ. ID NO.:434)  | HCV NS3 579-587  | 74 |
| TLGIVCPIC<br>(SEQ. ID NO.:435)  | HPV16 E7 86-94   | 74 |
| YLLPRRGPRL<br>(SEQ. ID NO.:436) | HCV core protein 34-43                                 | 74 |
| LLPIFFCLWV<br>(SEQ. ID NO.:437) | HBV env 378-387  | 74 |
| YMDDVVLGA<br>(SEQ. ID NO.:438)  | HBV Pol 538-546  | 74 |
| GTLGIVCPI<br>(SEQ. ID NO.:439)  | HPV16 E7 85-93   | 74 |
| LLALLSCLTI<br>(SEQ. ID NO.:440) | HCV MP 63-72   | 74 |

**TABLE II**  
**HLA Class I Motifs**

|                                 |                                     |    |
|---------------------------------|-------------------------------------|----|
| MLDLQPETT<br>(SEQ. ID NO.:441)  | HPV16 E7 12-20                      | 74 |
| SLMAFTAAV<br>(SEQ. ID NO.:442)  | HCV NS4 174-182                     | 75 |
| CINGVCWTV<br>(SEQ. ID NO.:443)  | HCV NS3 67-75                       | 75 |
| VMNILLQYVV<br>(SEQ. ID NO.:444) | Glutamic acid decarboxylase 114-123 | 76 |
| ILTVILGVL<br>(SEQ. ID NO.:445)  | Melan A/Mart- 32-40                 | 77 |
| FLWGPRAV<br>(SEQ. ID NO.:446)   | MAGE-3 271-279                      | 78 |
| LLCPAGHAV<br>(SEQ. ID NO.:447)  | HCV NS3 163-171                     | 54 |
| ILDSFDPLV<br>(SEQ. ID NO.:448)  | HCV NSS 239-247                     | 54 |
| LLLCLIFLL<br>(SEQ. ID NO.:449)  | HBV env 250-258                     | 79 |
| LIDYQGMLPV<br>(SEQ. ID NO.:450) | HBV env 260-269                     | 79 |
| SIVSPFIPLL<br>(SEQ. ID NO.:451) | HBV env 370-379                     | 79 |
| FLLTRILTI<br>(SEQ. ID NO.:452)  | HBV env 183-191                     | 80 |
| HLGNVKYLV<br>(SEQ. ID NO.:453)  | P. faciparum TRAP 3-11              | 81 |
| GIAGGLALL<br>(SEQ. ID NO.:454)  | P. faciparum TRAP 500-508           | 81 |
| ILAGYGAGV<br>(SEQ. ID NO.:455)  | HCV NS S4A 236-244                  | 82 |
| GLQDCTMLV<br>(SEQ. ID NO.:456)  | HCV NS5 714-722                     | 82 |
| TGAPVTYSTY<br>(SEQ. ID NO.:457) | HCV NS3 281-290                     | 83 |
| VIYQYMDDL<br>(SEQ. ID NO.:458)  | HIV-1RT 179-187                     | 84 |

**TABLE II**  
**HLA Class I Motifs**

|                                   |   |    |
|-----------------------------------|---|----|
| VLPDV FIRC V<br>(SEQ. ID NO.:459) | N-acetylglucosaminyltransferase V<br>Gnt-V intron | 85 |
| VLPDV FIRC<br>(SEQ. ID NO.:460)   | N-acetylglucosaminyltransferase V<br>Gnt-V intron | 85 |
| AVGIGIA V V<br>(SEQ. ID NO.:461)  | Human CD9   | 86 |
| LVVLG LLA V<br>(SEQ. ID NO.:462)  | Human glutamyltransferase                         | 86 |
| ALGLG LLP V<br>(SEQ. ID NO.:463)  | Human G protein coupled receptor<br>164-172       | 86 |
| GIGIGV LAA<br>(SEQ. ID NO.:281)   | HSV-1 gp C 480-488                                | 86 |
| GAGIGV AVL<br>(SEQ. ID NO.:464)   | HSV-2 gp C 446-454                                | 86 |
| IAGIGILAI<br>(SEQ. ID NO.:465)    | Pseudorabies gpGIN 455-463                        | 86 |
| LIVIGILIL<br>(SEQ. ID NO.:466)    | Adenovirus 3 E3 9kD 30-38                         | 86 |
| LAGIGLIAA<br>(SEQ. ID NO.:467)    | S. Lincolnensis ImrA                              | 86 |
| VDGIGILT I<br>(SEQ. ID NO.:468)   | Yeast ysa-1 77-85                                 | 86 |
| GAGIGV LTA<br>(SEQ. ID NO.:469)   | B. polymyxa, $\beta$ -endoxylanase 149-157        | 86 |
| AAGIGIIQ I<br>(SEQ. ID NO.:470)   | E. coli methionine synthase 590-598               | 86 |
| QAGIGILLA<br>(SEQ. ID NO.:471)    | E. coli hypothetical protein 4-12                 | 86 |
| KARDPHSGHFV<br>(SEQ. ID NO.:472)  | CDK4 <sup>wl</sup> 22-32                          | 87 |
| KACDPHSGHFV<br>(SEQ. ID NO.:473)  | CDK4-R24C 22-32                                   | 87 |
| ACDPHSGHFV<br>(SEQ. ID NO.:474)   | CDK4-R24C 23-32                                   | 87 |
| SLYNTVATL<br>(SEQ. ID NO.:475)    | HIV-1 gag p17 77-85                               | 88 |

**TABLE II**  
**HLA Class I Motifs**

|                                 |                                |    |
|---------------------------------|--------------------------------|----|
| ELVSEFSRV<br>(SEQ. ID NO.:476)  | HER-2, m>V substituted 971-979 | 89 |
| RGPGRFVTI<br>(SEQ. ID NO.:477)  | HIV-1 gp160 315-329            | 90 |
| HMWNFISGI<br>(SEQ. ID NO.:478)  | HCV NS4A 149-157               | 91 |
| NLVPMVATVQ<br>(SEQ. ID NO.:479) | HCMV pp65 495-504              | 92 |
| GLHCYEQLV<br>(SEQ. ID NO.:480)  | HPV 6b E7 21-30                | 93 |
| PLKQHFQIV<br>(SEQ. ID NO.:481)  | HPV 6b E7 47-55                | 93 |
| LLDFVRFMGV<br>(SEQ. ID NO.:482) | EBNA-6 284-293                 | 95 |
| AIMEKNIML<br>(SEQ. ID NO.:483)  | Influenza Alaska NS 1 122-130  | 67 |
| YLKTIQNSL<br>(SEQ. ID NO.:484)  | P. falciparum cp36 CSP         | 96 |
| YLNKIQNSL<br>(SEQ. ID NO.:485)  | P. falciparum cp39 CSP         | 96 |
| YMLDLQPETT<br>(SEQ. ID NO.:486) | HPV16 E7 11-20***              | 97 |
| LLMGTLGIV<br>(SEQ. ID NO.:487)  | HPV16 E7 82-90 ***             | 97 |
| TLGIVCPI<br>(SEQ. ID NO.:488)   | HPV16 E7 86-93***              | 97 |
| TLTSCNTSV<br>(SEQ. ID NO.:489)  | HIV-1 gp120 197-205            | 98 |
| KLPQLCTEL<br>(SEQ. ID NO.:490)  | HPV16 E6 18-26                 | 97 |
| TIHDIILEC<br>(SEQ. ID NO.:491)  | HPV16 E6 29-37                 | 97 |
| LGIVCPICS<br>(SEQ. ID NO.:492)  | HPV16 E7 87-95                 | 97 |
| VILGVLLLI<br>(SEQ. ID NO.:493)  | Melan A/Mart-1 35-43           | 68 |

**TABLE II**  
**HLA Class I Motifs**

|                 |                                 |                               |     |
|-----------------|---------------------------------|-------------------------------|-----|
|                 | ALMDKSLHV<br>(SEQ. ID NO.:494)  | Melan A/Mart-1 56-64          | 68  |
|                 | GILTVILGV<br>(SEQ. ID NO.:495)  | Melan A/Mart-1 31-39          | 68  |
| T cell epitopes | MINAYLDKL<br>(SEQ. ID NO.:496)  | P. Falciparum STARP 523-531   | 81  |
|                 | AAGIGILTV<br>(SEQ. ID NO.:497)  | Melan A/Mart-1 27-35          | 100 |
|                 | FLPSDFFPSV<br>(SEQ. ID NO.:498) | HBV cAg 18-27                 | 51  |
| Motif unknown   | SVRDRLARL<br>(SEQ. ID NO.:499)  | EBNA-3 464-472                | 101 |
| T cell epitope  |                                 |                               |     |
| T cell epitope  | AAGIGILTV<br>(SEQ. ID NO.:497)  | Melan A/Mart-1 27-35          | 100 |
|                 | FAYDGKDYI<br>(SEQ. ID NO.:500)  | Human MHC 1- $\alpha$ 140-148 | 99  |
| T cell epitopes | AAGIGILTV<br>(SEQ. ID NO.:497)  | Melan A/Mart-1 27-35          | 100 |
|                 | FLPSDFFPSV<br>(SEQ. ID NO.:498) | HBV cAg 18-27                 | 51  |
|                 | AAGIGILTV<br>(SEQ. ID NO.:497)  | Melan A/Mart-1 27-35          | 100 |
|                 | FLPSDFFPSV<br>(SEQ. ID NO.:498) | HBV cAg 18-27                 | 51  |
| Motif unknown   | AAGIGILTV<br>(SEQ. ID NO.:497)  | Melan A/Mart-1 27-35          | 100 |
| T cell epitope  |                                 |                               |     |
|                 | ALLAVGATK<br>(SEQ. ID NO.:501)  | Pmel17 gp100 17-25            | 107 |
| T cell epitopes | RLRDLLIVTR<br>(SEQ. ID NO.:502) | HIV-1 gp41 768-778            | 108 |
|                 | QVPLRPMTYK<br>(SEQ. ID NO.:503) | HIV-1 nef 73-82               | 109 |
|                 | TVYYGVVWK<br>(SEQ. ID NO.:504)  | HIV-1 gp120-36-45             | 110 |
|                 | RLRPGGKKK<br>(SEQ. ID NO.:505)  | HIV-1 gag p17 20-29           | 110 |

**TABLE II**  
**HLA Class I Motifs**

|                                 |                                     |  |                               |
|---------------------------------|-------------------------------------|--|-------------------------------|
|                                 | ILRGSAHK<br>(SEQ. ID NO.:506)       | Influenza NP 265-273                                 | 21                            |
|                                 | RLRAEAGVK<br>(SEQ. ID NO.:507)      | EBNA-3 603-611                                       | 111                           |
|                                 | RLRDLLIVTR<br>(SEQ. ID NO.:502)     | HIV-1 gp41 770-780                                   | 112                           |
|                                 | VYYGVVPWK<br>(SEQ. ID NO.:508)      | HIV-1 GP120 38-46                                    | 113                           |
|                                 | RVCEKMALY<br>(SEQ. ID NO.:509)      | HCV NS5 575-583                                      | 114                           |
| Motif unknown<br>T cell epitope | KIFSEVTLK<br>(SEQ. ID NO.:510)      | Unknown; mutated (p183L)<br>melanoma peptide 175-183 | Wolfel et al., pers.<br>Comm. |
|                                 | YVNVNMG LK*<br>(SEQ. ID NO.:511)    | HBV cAg 88-96  | 116                           |
| T cell epitopes                 | IVTDFSVIK<br>(SEQ. ID NO.:512)      | EBNA-4 416-424                                       | 115, 117                      |
|                                 | ELNEALELK<br>(SEQ. ID NO.:513)      | P53 343-351  | 115                           |
|                                 | VPLRPMTYK<br>(SEQ. ID NO.:514)      | HIV-1 NEF 74-82                                      | 115                           |
|                                 | AIFQSSMTK<br>(SEQ. ID NO.:515)      | HIV-1 gag p24 325-333                                | 115                           |
|                                 | QVPLRPMTYK<br>(SEQ. ID NO.:516)     | HIV -1 nef 73-82                                     | 118                           |
|                                 | TINYTIK<br>(SEQ. ID NO.:517)        | HCV NSI 238-246                                      | 114                           |
|                                 | AAVDLSHFLKEK<br>(SEQ. ID NO.:518)   | HIV-1 nef 83-94                                      | 120                           |
|                                 | ACQGVGGPGGHK<br>(SEQ. ID NO.:519)   | HIV-1 111B p24 349-359                               | 122                           |
| HLA-A24                         | SYLD SG I HF*<br>(SEQ. ID NO.:520)  | $\beta$ -catenin, mutated (proto-onocogen)<br>29-37  | 123                           |
| T cell epitopes                 | RYLK D Q Q L L<br>(SEQ. ID NO.:521) | HIV GP 41 583-591                                    | 124                           |
|                                 | AYGLDFYIL<br>(SEQ. ID NO.:522)      | P15 melanoma Ag 10-18                                | 125                           |

**TABLE II**  
**HLA Class I Motifs**

|                 |  |                               |          |
|-----------------|--|-------------------------------|----------|
|                 | A F L P W H R L F L<br>(SEQ. ID NO.:523)   | Tyrosinase 206-215            | 126      |
|                 | A F L P W H R L F<br>(SEQ. ID NO.:524)     | Tyrosinase 206-214            | 126      |
|                 | R Y S I F F D Y<br>(SEQ. ID NO.:525)       | Ebna-3 246-253                | 101      |
| T cell epitope  | E T I N E E A A E W<br>(SEQ. ID NO.:526)   | HIV-1 gag p24 203-212         | 127      |
| T cell epitopes | S T I P E T T V V R R<br>(SEQ. ID NO.:527) | HBV cAg 141-151               | 129      |
|                 | M S L Q R Q F L R<br>(SEQ. ID NO.:528)     | ORF 3P-gp75 294-321 (bp)      | 130      |
|                 | I L P G G R P Y R<br>(SEQ. ID NO.:528)     | TRP (tyrosinase rel.) 197-205 | 131      |
| T cell epitope  | I V G L N K I V R<br>(SEQ. ID NO.:530)     | HIV gag p24 267-267-275       | 132, 133 |
|                 | A A G I G I L T V<br>(SEQ. ID NO.:531)     | Melan A/Mart-1 27 35          | 100      |

Table III sets forth additional antigens useful in the invention that are available from the Ludwig Cancer Institute. The Table refers to patents in which the identified antigens can be found and as such are incorporated herein by reference. TRA refers to the tumor-related antigen and the LUD No. refers to the Ludwig Institute number.

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| Table III  |         |            |                    |  |         |
|------------|---------|------------|--------------------|--|---------|
| TRA        | LUD No. | Patent No. | Date Patent Issued | Peptide (Antigen)  | HLA     |
| MAGE-4     | 5293    | 5,405,940  | 11 April 1995      | EVDPASNTY<br>(SEQ. ID NO.:532)                                 | HLA-A1  |
| MAGE-41    | 5293    | 5,405,940  | 11 April 1995      | EVDPTSNTY<br>(SEQ ID NO:533)                                   | HLA-A1  |
| MAGE-5     | 5293    | 5,405,940  | 11 April 1995      | EADPTSNTY<br>(SEQ ID NO:534)                                   | HLA-A1  |
| MAGE-51    | 5293    | 5,405,940  | 11 April 1995      | EADPTSNTY<br>(SEQ ID NO:534)                                   | HLA-A1  |
| MAGE-6     | 5294    | 5,405,940  | 11 April 1995      | EVDPIGHVY<br>(SEQ ID NO:535)                                   | HLA-A1  |
|            | 5299.2  | 5,487,974  | 30 January 1996    | MLLAVLYCLL<br>(SEQ ID NO:536)                                  | HLA-A2  |
|            | 5360    | 5,530,096  | 25 June 1996       | MLLAVLYCL<br>(SEQ ID NO:537)                                   | HLA-B44 |
| Tyrosinase | 5360.1  | 5,519,117  | 21 May 1996        | SEIWRDIDFA<br>(SEQ ID NO:538)<br>SEIWRDIDF<br>(SEQ ID NO:539)  | HLA-B44 |
| Tyrosinase | 5431    | 5,774,316  | 28 April 1998      | XEIWRDIDF<br>(SEQ ID NO:540)                                   | HLA-B44 |
| MAGE-2     | 5340    | 5,554,724  | 10 September 1996  | STLVEVTLGEV<br>(SEQ ID NO:541)<br>LVEVTLGEV<br>(SEQ ID NO:542) | HLA-A2  |



Table III

| TRA         | LUD<br>No. | Patent<br>No. | Date Patent<br>Issued | Peptide (Antigen)  | HLA               |
|-------------|------------|---------------|-----------------------|--|-------------------|
| (Continued) |            |               |                       | VIFSKASEYL<br>(SEQ ID NO:543)  |                   |
|             |            |               |                       | IIVLAIIAI<br>(SEQ ID NO:544)   |                   |
|             |            |               |                       | KIWEELSMLEV<br>(SEQ ID NO:545)   |                   |
|             |            |               |                       | LIETSYVKV<br>(SEQ ID NO:546)   |                   |
|             |            |               |                       |  |                   |
|             |            |               |                       |  |                   |
|             | 5327       | 5,585,461     | 17 December 1996      | FLWGPRALV<br>(SEQ ID NO:547)   | HLA-A2            |
|             |            |               |                       | TLVEVTLGEV<br>(SEQ ID NO:548)  |                   |
|             |            |               |                       | ALVETSYVKV<br>(SEQ ID NO:549)  |                   |
| MAGE-3      | 5344       | 5,554,506     | 10 September 1996     | KIWEELSVL<br>(SEQ ID NO:550)   | HLA-A2            |
| MAGE-3      | 5393       | 5,405,940     | 11 April 1995         | EVDPIGHLV<br>(SEQ ID NO:551)   | HLA-A1            |
| MAGE        | 5293       | 5,405,940     | 11 April 1995         | EXDX <sub>5</sub> Y<br>(SEQ. ID NO.:552)<br>(but not EADPTGHSY)<br>(SEQ. ID NO.:553)<br>E (A/V) D X <sub>5</sub> Y<br>(SEQ. ID NO.:554)<br>E (A/V) D P X <sub>4</sub> Y<br>(SEQ. ID NO.:555)<br>E (A/V) D P (I/A/T) X <sub>3</sub> Y<br>(SEQ. ID NO.:556)<br>E (A/V) D P (I/A/T) (G/S) X <sub>2</sub> Y<br>(SEQ. ID NO.:557)<br>E (A/V) D P (I/A/T) (G/S) (H/N) X Y<br>(SEQ. ID NO.:558)<br>E (A/V) D P (I/A/T) (G/S) (H/N) (L/T/V) Y<br>(SEQ. ID NO.:559) | HLA-A1            |
| MAGE-1      | 5361       | 5,558,995     | 24 September 1996     | ELHSAYGEPRKLLTQD<br>(SEQ ID NO:560)  | HLA-C<br>Clone 10 |

Table III

| TRA    | LUD<br>No. | Patent<br>No. | Date Patent<br>Issued | Peptide (Antigen)   | HLA   |
|--------|------------|---------------|-----------------------|---|---|
|        |            |               |                       | EHSAYGEPRKLL<br>(SEQ ID NO:561)<br>SAYGEPRKL<br>(SEQ ID NO:562)   |   |
| MAGE-I | 5253.4     | TBA           | TBA                   | EADPTGHSY<br>(SEQ ID NO:563)  | HLA-A1  |
| BAGE   | 5310.1     | TBA           | TBA                   | MAARAVFLALSAQILLQARLMKE<br>(SEQ ID NO:564)<br>MAARAVFLALSAQLLQ<br>(SEQ ID NO:565)<br>AARAVFLAL<br>(SEQ ID NO:566) | HLA-C<br>Clone 10<br>HLA-C<br>Clone 10<br>HLA-C<br>Clone 10 |
| GAGE   | 5323.2     | 5,648,226     | 15 July 1997          | YRPRPRRY<br>(SEQ. ID NO.:567)   | HLA-CW6   |

Preferred peptide antigens are those of tumor associated antigens (TAA) and chronic infections. Particularly preferred peptide antigens are derived from tyrosinase, gp100 or Melan A for the treatment of melanoma.

5 The peptide antigens of this invention are readily prepared using standard peptide synthesis means known in the art. Generally they can be prepared commercially by one of numerous companies that do chemical synthesis. An example is American Peptides, Inc., where the distributor is CLINALFA AG (Laufelfingen, Switzerland). The antigens are prepared in accordance with GMP standards. Purity is assessed by analytical HPLC. The product is characterized by  
10 amino-acid analysis and tested for sterility and the absence of pyrogens.

In delivering an appropriate antigen, *e.g.*, a polypeptide, to the animal's system it may be delivered directly as the polypeptide, or it may be delivered indirectly, *e.g.*, using a DNA construct or vector, or a recombinant virus that codes for the desired antigen. Any vector driving expression in a professional antigen  
15 presenting cell is suitable for this purpose. In the indirect delivery, the antigen is expressed in the cell, to be presented by the MHC Class I on the surface of the cell to stimulate the CTL response.

Antigens may be used alone or may be delivered in combination with other antigens or with other compounds such as cytokines that are known to enhance  
20 immune stimulation of CTL responses, such as, GM-CSF, IL-12, IL-2, TNF, IFN, IL-18, IL-3, IL-4, IL-8, IL-9, IL-13, IL-10, IL-14, IL-15, G-CSF, IFN alpha, IFN beta, IFN gamma, TGF alpha, TGF beta, and the like. The cytokines are known in the art and are readily available in the literature or commercially. Many animal and human tumors have been shown to produce cytokines such as IL-4, IL-10, TGF- $\beta$  that are  
25 potent modulators of the immune response and that protect tumors from immune-mediated destruction. The production of IL-4, IL-10 or TGF- $\beta$  by the tumors may achieve this protective effect by suppressing the induction of cellular immunity, including the elaboration of CTL responses. Alternatively, cytokines that support CTL responses can be exogenously added to help in the balance between induction of  
30 anti-tumor cell mediated and non-tumor-destructive humoral responses. Several such exogenous cytokines show utility in experimental mouse vaccination models which are known to enhance CTL responses, including GM-CSF, IFN and IL-2. An effective exogenous cytokine that may be used is GM-CSF. GM-CSF is reported to enhance the expression of the so called "co-stimulatory" molecules, such as B7-1 or

B7-2 on antigen presenting cells (APC), which are important players in the symphony of interactions that occur during stimulation of CTL by APC. Moreover, GM-CSF is known to induce activation of APC and to facilitate growth and differentiation of APC, thereby making these important CTL stimulating cells available both in greater  
5 numbers and potency.

### **Delivery of the Antigen**

This invention is based in part on the observation that a CTL response is not sustained using standard vaccine techniques. While not wanting to be bound by any particular theory, it is thought that T cells do not have a functional memory that is  
10 long-lived. Antibody-mediated B-cell memory, on the other hand, appears to have a long-lived effector memory. Thus, delivering an antigen that produces a CTL response must be done over time to keep the patient's immune system appropriately stimulated to attack the target cells. While it has been suggested that antigens and  
15 adjuvants can be prepared as biodegradable microspheres or liposomes, none of these preparations have thus far provided a CTL response that is useful for attacking cancer cells or pathogens on a long term basis. The delivery must be sustained over the desired period of time at a level sufficient to maintain the antigen level to obtain the desired response and that it must be delivered from a reservoir having fluid antigen  
composition that is introduced so that it reaches the animal's lymphatic system.

20 Ultimately antigen must find its way into the lymphatic system in order to efficiently stimulate CTL. However, delivery of antigen according to the invention can involve infusion into various compartments of the body, including but not limited to subcutaneous, intravenous, intraperitoneal and intralymphatic, the latter being preferred. Each of these various points of infusion results in antigen uptake into the  
25 lymphatic system. The relative amounts of antigen needed to induce a beneficial CTL response varies according to the different sites of infusion. In general, direct infusion of antigen into the lymph system is deemed to be the most efficient means of inducing a CTL response, but that the material difference between the various routes is not necessarily relevant in terms of the quantity of antigen needed, or, in terms of the  
30 operating parameters of the invention. The pump systems of the invention are capable of delivering material quantities of antigen in a range that makes the invention suitable for inducing CTL response through delivery to all compartments of the body. CTL stimulation based on delivery of antigen via various routes will be variable, based on the properties of different antigens, which will reflect factors that influence

antigen behavior in the body and its rate of equilibration to (or longevity in) the lymph, such as antigen stability in the body fluid, solubility of antigen in body fluid, binding affinity for HLA and potency as a stimulator of CTL.

It is most efficient, and therefore, preferred, that the introduction is done as directly as possible to the lymphatic system to avoid the destruction of the antigen by metabolism in the body. When introduction of a fluid antigen composition occurs subcutaneously, larger quantities of antigen are needed to assure enough antigen reaches the lymphatic system. Such subcutaneous injection is contemplated by this invention if it can be justified by factors such as cost, stability of the antigen, how quickly the antigen gets to the lymph system, how well it equilibrates with the lymph, and other factors that the attending doctor or specialist will recognize. Subcutaneous delivery will generally require 100 to 1000 times more antigen than direct delivery to the lymph system. It is preferable, therefore, that the antigen composition is introduced through a device for local administration to the lymphatic system, e.g. the spleen, a lymph node, or a lymph vessel. The device for local administration may be positioned outside the patient or implanted into the patient. In either case, the device will have a reservoir to hold the fluid antigen-containing composition, a pump to transfer the composition, and a transmission channel leading from the reservoir to be directed to the preferred region of administration in the patient's body. In either case it is preferably portable.

For the device positioned outside the patient's body (the external device), there are numerous devices used for delivering insulin to diabetic patients that are useful in this invention. Generally these are comprised of a reservoir for holding the antigen composition (instead of insulin), a programmable pump to pump the composition out of the reservoir, a transmission channel or line for transmitting the composition, and a means to introduce the composition into the animal's body to ultimately reach the lymphatic system.

The pump employed may be a roller/peristaltic pump, a syringe pump, a piston/valve pump, a gas pressure pump, or the like that has a power source (generally a battery for portability) that is programmable to deliver the desired level of antigen composition to the patient's body and the lymphatic system. A further discussion of the operation of these pumps may be found "Insulin Pump Therapy" by E. Austenst and T. Stahl, Walter de Gruyter, Berlin, New York (1990), at Chapter 3. A list of

pumps available at that time that are useful for this invention are given in Table IV. More recent versions of these pumps are available from the manufacturers shown.

TABLE IV

| Name                   | Manufacturer/distributor       | Weight (g) | Size (mm)     |
|------------------------|--------------------------------|------------|---------------|
| Nordisk Infusor        | Nordisk                        | 180        | 100 x 60 x 20 |
| Betatron I             | CPI/Lilly                      | 197        | 99 x 66 x 20  |
| RW 90 P/RW 91 P/ RW 92 | Dahedi/EA Satorius Instruments | 110        | 109 x 42 x 22 |
| MRS 4-Infuser          | Disetronic                     | 100        | 75 x 53 x 18  |
| B-D 1000               | Becton-Dickinson               | 131        | 78 x 57 x 20  |
| Nordisk Infusor MK II  | Nordisk                        | 180        | 113 x 65 x 22 |
| MRS 3-Infuser          | Disetronic                     | 100        | 75 x 53 x 18  |
| A S8 MP                | Autosyringe/Travenol           | 161        | 102 x 64 x 19 |
| Betatron II            | CPI/Lilly                      | 197        | 99 x 66 x 20  |
| Minimed 504            | Pacesetter/Haselmeyer          | 106        | 86 x 21 x 51  |
| Minimed 404 S*         | Pacesetter                     | 106        | 86 x 21 x 51  |
| MRS 1/H-Tron           | Disetronic/Hoechst             | 100        | 75 x 53 x 18  |

\* not yet commercially available

- 5 Particularly useful pumps are the Disetronic H-Tron V100 Insulin Pump from Disetronic Medical Systems, Burgdorf, Switzerland and the Minimed 507 Insulin Pump from MiniMed Inc., 12744 San Fernando Road, Sylmar, California 91342. The MiniMed is particularly useful in that it allows programming a bolus without looking at the pump through a series of audio tones (settable in either 0.5 or 1.0 unit
- 10 increments) and allows programming a bolus for delivery over an extended period of time – from 30 minutes to 4 hours. It provides up to 12 basal rates (or profiles) that can be programmed per 24 hours from 0.0 –25 units/hour in 0.1 unit increments. The device allows for the temporary increase or decrease of a set basal rate from 30 minutes to 24 hours in 30 minute increments. Other features relating to safety, time
- 15 display, memory, etc. are available from the manufacturer.

The reservoir for the antigen composition should be large enough for delivery of the desired amount of antigen over time and is easily refillable or replaceable

without requiring the user to reinsert the means for introducing the antigen composition to the lymph system.

In preparing the antigen compositions of this invention, a composition (preferably aqueous) is prepared to be compatible with the lymph system and is physiologically acceptable to the animal being treated. In preparing the antigen compositions useful in this invention one considers the physicochemical properties of the antigen such as the isoelectric point, molecular weight, glycosylation or other post-translational modification, and overall amino acid composition. These properties along with any known behavior of the drug in different solutions (e.g. different buffers, cofactors, etc.) as well as its *in vivo* behavior will help guide the choice of formulation components. One parameter that impacts all the major degradation pathways is the solution pH. Thus, the initial formulations also assess the pH dependence of the degradation reactions and the mechanism for degradation can often be determined from the pH dependence to determine the stability of the protein in each solution. Rapid screening methods usually involve the use of accelerated stability at elevated temperatures (e.g. 40° C) using techniques known in the art.

In general the antigen compositions useful in this invention will be prepared suitable for parenteral injection, in very small quantities. As such a composition must be free of contamination and have a pH compatible with the lymph system. However, because very small quantities of the antigenic composition will be delivered it need not be the same pH as blood or lymph, and it need not be aqueous-based. For antigens that are less soluble a suitable cosolvent or surfactant may be used, such as dimethyl sulfoxide (DMSO) or PLURONIC brand surfactants. The pH range that is compatible is from about 6.7 – 7.3 and can be prepared using water for injection to meet USP specifications (see Remington: The Science and Practice of Pharmacy, Nineteenth Edition; Chapters 86-88). Generally, a standard saline solution that is buffered with a physiologically acceptable weak acid and its base conjugate, e.g., a phosphate or citrate buffering system, will be the basis of the antigen composition. In some cases, a small amount of an antioxidant may be useful to stabilize the composition and prevent oxidation. Factors to consider in preparing the antigen compositions may be found in the 1994 American Chemical Society book entitled "*Formulation and Delivery of Proteins and Peptides*" (Acs Symposium Series, No. 567) by Jeffery L. Cleland and Robert Langer (Editor)).

Generally the amount of the antigen in the antigen composition will vary from patient to patient and from antigen to antigen, depending on such factors as the activity of the antigen in inducing a response and the flow rate of the lymph through the patient's system. In general the antigen composition may be delivered at a rate of  
5 from about 1 to about 500 microliters/hour or about 24 to about 12000 microliters/day. The concentration of the antigen is such that about 0.1 micrograms to about 10,000 micrograms of the antigen will be delivered during 24 hours. The flow rate is based on the knowledge that each minute approximately about 100 to about 1000 microliters of lymph fluid flows through an adult inguinal lymph node. The  
10 objective is to maximize local concentration of vaccine formulation in the lymph system. A certain amount of empirical investigation on patients will be necessary to determine the most efficacious level of infusion for a given vaccine preparation in humans.

To introduce the antigen composition into the lymphatic system of the patient,  
15 the composition is preferably directed to a lymph vessel, lymph node, the spleen, or other appropriate portion of the lymph system. Preferably, the composition is directed to a lymph node such as an inguinal or axillary node by inserting a catheter or needle to the node and maintaining the catheter or needle throughout the delivery. Suitable needles or catheters are available made of metal or plastic (e.g. polyurethane,  
20 polyvinyl chloride [PVC], TEFLON, polyethylene, and the like). In inserting the catheter or needle into the inguinal node for example, the inguinal node is punctured under ultrasonographic control using a Vialon™ Insyte-W™ cannula and catheter of 24G<sub>3/4</sub> (Becton Dickinson, USA) which is fixed using Tegaderm transparent dressing (Tegaderm™ 1624, 3M, St. Paul, MN 55144, USA). This procedure is generally  
25 done by an experienced radiologist. The location of the catheter tip inside the inguinal lymph node is confirmed by injection of a minimal volume of saline, which immediately and visibly increases the size of the lymph node. The latter procedure allows confirmation that the tip is inside the node and can be performed to ensure that the tip does not slip out of the lymph node can be repeated on various days after  
30 implantation of the catheter. In case the tip did in fact slip out of location inside the lymph node, a new catheter can be implanted.

In another embodiment, the antigen is delivered to the lymphatic system through an article of manufacture that is implanted in the animal, preferably at or near a site of a lymphatic organ. The article will include a pump that can deliver the



antigen at a controlled rate over a pre-determined period of time and is suitable for use in the host. Several devices are known in the art for the delivery of agents (such as drugs) in humans or animals and these can be used or adapted for use in the present invention.

5           The implantable device will be similar to the external device discussed above in that it comprises a reservoir of a physiologically-acceptable, aqueous, antigen-containing composition that is capable of inducing a CTL response in an animal, a pump positioned in association with the reservoir to deliver the composition at a defined rate, a transmission channel to discharge the composition from the reservoir,  
10           and optionally a delivery line connected to the transmission channel, which delivery line is of a size suitable for positioning in the animal and for delivery of the composition in a manner that reaches the lymphatic system of the animal.

          Preferably the pump in the implantable device is an osmotic pump of the type used in the ALZET® model device or the DUROS™ model device pioneered by Alza  
15           Corporation, Palo Alto, CA or in a device made by Pharmetrix and exemplified in U.S. patent 4,838,862. The osmotic pump utilizes the osmotic effect using a membrane permeable to water but impermeable to a solute. Osmotic pressure built up in a device is used to deliver a composition at a controlled rate over time. A review by Giancarlo Santus and Richard Baker of "Osmotic Drug Delivery: A Review of the  
20           Patent Literature" in the Journal of Controlled Release 35 (1995) 1-21, provides useful guidelines for the type of osmotic pumps that are useful in this invention. The osmotic pump forces the composition through a discharge orifice to discharge the composition. Optionally a delivery line connects to the discharge orifice to position the line suitably for delivery to the lymphatic system of the animal. Patents that  
25           describe devices useful in this invention include the following U.S. patents: (A) 3,604,417 assigned to American Cyanamid; (B) 4,838,862; 4,898,582; 5,135,498; 5,169,390; and 5,257,987 all assigned to Pharmetrix, (C) 4,340,048; 4,474,575; 4,552,651; 4,619,652; 4,753,651; 3,732,865; 3,760,804; 3,760,805; 3,929,132; 3,995,632; 4,034,756; 4,350,271; 4,455,145; 5,017,381; 5,023,088; 5,030,216;  
30           5,034,229; 5,037,420; 5,057,318; 5,059,423; 5,110,596; 5,110,597; 5,135,523; 5,137,727; 5,174,999; 5,209,746; 5,221,278; 5,223,265; 3,760,984; 3,987,790; 3,995,631; 4,203,440; 4,286,067; 4,300,558; 4,304,232; 4,340,054; 4,367,741; 4,450,198; 4,855,141; 4,865,598; 4,865,845; 4,872,873; 4,929,233; 4,963,141;

4,976,966, all assigned to Alza Corp. Each of the foregoing patents are incorporated herein by reference.

5 A basic osmotic pump device incorporates a housing containing a chamber for storing the antigen containing composition to be delivered, separated from a compartment containing an osmotic salt material by a barrier that is moveable under pressure such as a piston or a flexible impermeable membrane. The compartment containing the osmotic salt is separated from osmotic fluid by a semipermeable membrane. In some embodiments, a fluid barrier, such as a foil sheet, isolates the osmotic salt chamber from the osmotic fluid, keeping the pump inactivated until  
10 removal of the barrier immediately prior to use. Other osmotic pump devices use body fluid as the osmotic fluid. In these devices a semipermeable membrane separates the osmotic salt compartment from body fluids, and the pump is activated once inserted into the body under exposure to body fluids. In either case, volumetric expansion of the osmotic salt compartment drives the expulsion of the stored antigen  
15 from the compartment and into the surrounding environment of the body. These pumps have been highly successful at achieving steady-state pumping and delivery of agents. The pumps are of a small size that can be inserted into a patient, with flexible consideration as to location. This is important in the case of CTL vaccines, since the inventor has determined that efficient induction of CTL responses is contingent on the  
20 antigen or antigen expression system being delivered into the lymphatic system, in order to ultimately achieve antigen delivery into a lymphatic organ such as the spleen. Antigen delivered into a lymph node is 100-1000 times more efficient at inducing CTL responses compared with conventional subcutaneous delivery. A modification to the osmotic pump incorporates a microcatheter attachment (i.e., the optional  
25 delivery line) at its discharge end, such that when the pump is implanted proximal to a lymphatic organ, such as a lymph node, the catheter can be inserted into the organ to facilitate delivery of the vaccine directly into the lymphatic system.

Prior to the administration of the antigen using any of the above vehicles, methods may be used to assist in the determination of the optimum location for the  
30 antigen delivery. For example, when using the osmotic pump, radiography may be used to image a patient's lymphatic flow, to determine where relatively high lymphatic drainage occurs, in order to decide upon an insertion position for the osmotic pump that maximizes delivery into the lymphatic system. Since each patient has unique lymphatic drainage profiles, imaging would be conducted for each

individual prior to insertion of osmotic pump for delivery of antigen. When using direct cannulation of the lymphatic vessel, such as in the use of osmotic or insulin pumps to deliver antigen, ultrasound can be used to position the needle directly into the lymphatic vessel and to monitor its positioning over the period of treatment.

5           The following non-limiting examples are illustrative of the present invention.

## EXAMPLES

### Materials and Methods For Examples 1-5

**Mice:** The generation of T cell receptor transgenic mice (TCR+ mice) in which approx. 90% of the CD8+ T cells express a TCR recognizing the immunodominant  
10 LCMV-glycoprotein epitope (gp-peptide aa33-41, p33:KAVYNFATC-SEQ ID NO:569) presented on H-2D<sup>b</sup>, has been described in detail. All experimental mice were between 8 and 12 weeks of age and bred and held under strict pathogen free conditions at the Institut Für Labortierkunde at the University of Zurich.

**Viruses:** LCMV (Armstrong strain) was originally obtained from Dr. M.B.A.  
15 Oldstone, Scripps Clinics and Research Foundation, LaJolla, San Diego, CA. Seed virus was grown on BHK cells and plaqued on MC57 cells using an immunological focus assay, as described previously.

**Osmotic pump:** ALZA model #1007b.

*In vivo* protection assays for specific CTL activity: The *in vivo* assay for the  
20 detection of CTL activity by challenge infections with LCMV has been described in detail previously (Oehen et al. 1991). Briefly, mice are intravenously challenged with  $2 \times 10^3$  pfu of LCMV (Armstrong). After 4 days the titer of LCMV is determined using the above mentioned immunological focus assay.

**Primary ex vivo cytotoxicity against LCMV-gp:** Mice were injected  
25 intravenously with 10µg of p33. After 36 hours spleen single cell suspensions were coincubated for 5h with <sup>51</sup>Cr-labeled syngeneic EL-4 (H-2<sup>b</sup>) target cells, that were either pulsed with p33 or left unpulsed. Specific lysis was calculated as  $[(\text{experimental } ^{51}\text{Cr release} - \text{spontaneous } ^{51}\text{Cr release}) / (\text{total } ^{51}\text{Cr release} - \text{spontaneous } ^{51}\text{Cr release}) \times 100\%]$ .

30           **LCMV induced foot pad swelling reaction:** Mice were infected with LCMV (Armstrong) by intradermal injection into the hind footpad (5000 pfu in 30:1). Footpad thickness was measured daily with a spring loaded caliper. Footpad swelling

is calculated as (measured thickness -thickness before injection) / (thickness before injection).

### Example 1

#### *Continuous release of peptide antigen using osmotic pump induces potent CTL response in C57BL/6 Mice*

C57BL/6 mice were either intravenously injected with a single dose of 50 $\mu$ g p33 (including 500 ng GM-CSF) (circles) or were implanted with a micro-osmotic pump releasing a mixture of 50 $\mu$ g of p33 and 500 ng GM-CSF over a time period of 7 days (triangles), or were left naïve (data not shown). After 7 days mice were sacrificed to prepare single cell suspensions from the spleen. Spleen cells were restimulated in vitro for 5 days by p33 pulsed in the presence of low amounts of IL-2. Specific cytotoxicity was measured using <sup>51</sup>Cr-labeled EL-4 target cells pulsed with p33. Specific lysis of EL-4 target cells without p33 was less than 16% for all effectors. The results are shown in Figure 1.

### Example 2

#### *Continuous release of antigen induces CTL immunity against virus in C57BL/6 mice*

C57BL/6 mice were either intravenously injected with a single dose of 50 $\mu$ g p33 (including 500 ng GM-CSF, Pharmingen) or were implanted with a micro-osmotic pump releasing a mixture of 50 $\mu$ g of p33 and 500 ng GM-CSF over a time period of 7 days, or were left naïve. After 7 days specific CTL activity was assessed *in vivo* using anti-viral protection assays. C57BL/6 mice were intravenously challenged with LCMV Armstrong strain (2x10<sup>3</sup> p.f.u.). After 4 days mice were sacrificed and LCMV titers were determined in spleens using an immunological focus assay. Mice implanted with osmotic pump showed significantly lower virus titers indicating active CTL immunity against the virus (Table V).

**TABLE V**

| <b>C57BL/6 Mice</b> | <b>Virus Titer (log<sub>10</sub>)</b> |
|---------------------|---------------------------------------|
| Single injection    | 4.2                                   |
| Single injection    | 4.6                                   |
| Single injection    | 4.0                                   |
| Pump delivered      | 2.2                                   |
| Pump delivered      | 1.8                                   |
| Pump delivered      | 2.0                                   |
| Unprimed            | 4.8                                   |
| Unprimed            | 3.8                                   |
| Unprimed            | 4.4                                   |

**Example 3****5 Continuous release of antigen maintains potent CTL effectors in TCR Transgenic Mice**

TCR transgenic mice were either intravenously injected with a single dose of 50µg p33 (circles) or were implanted with a microsomatic pump releasing a mixture of 50µg of p33 (triangles), or left naïve (squares). After 36 hours mice were sacrificed to prepare single cell suspensions from the spleen which were assayed ex vivo for p33-specific cytotoxicity using <sup>51</sup>Cr-labeled EL-4 target cells pulsed with p33. Similarly mice were either intravenously injected with a single dose of 50µg p33 (circles) or were implanted with a micro-osmotic pump releasing a mixture of 50µg of p33 over a time period of 7 days (triangles), or were left naïve (squares). After 7 days mice were sacrificed to prepare single cell suspensions from the spleen to assay ex vivo p33-specific cytotoxicity using <sup>51</sup>Cr-labeled EL-4 target cells pulsed with p33. Specific lysis of EL-4 target cells without p33 was less than 18% for all effectors. The results are shown in Figures 2A and 2B.

**20 Continuous release of antigen maintains protective CTL response against virus infection.**

After 7 days TCR transgenic mice were challenged by intradermal LCMV injection into their hind foot pads (2x10<sup>3</sup> pfu in 30µl). The absence of a foot pad swelling reaction, as observed in mice with an implanted pump (triangles), indicates that at the time point of injection there was active CTL immunity inhibiting local replication of the virus in the foot pad. In contrast, foot pad swelling, as observed in mice injected with the peptide as a single bolus (circles) and naïve control mice (data

not shown), indicated that LCMV successfully replicated in the foot pad in the absence of protective CTL. The results are shown in Figure 2C.

#### **Example 4**

##### ***Direct delivery of antigen into lymphatic organ dramatically increases efficiency of CTL induction***

TCR transgenic mice were injected with graded doses of gp-peptide p33 either subcutaneously (S.C.), intravenously (I.V.) or directly into the spleen (I.S.) via a small abdominal incision. The efficiency of CTL induction was assessed by measuring gp-specific CTL activity 24 hours after injection. CTL activity is known to peak one day after injection of peptide. Mice were sacrificed to prepare single cell suspensions from draining lymph nodes or from spleen to assay ex vivo p33-specific cytotoxicity using <sup>51</sup>Cr-labeled EL-4 target cells pulsed with p33. Specific lysis of EL-4 target cells without p33 was less than 12% for all effectors. The results are shown in Figure 3.

#### **Example 5**

##### ***Dendritic Cells Purified from Mice Receiving Intrasplenic Injection of Peptide Potently Stimulate CTL***

The effect of directing peptide delivery into lymphatic system was assessed. Peptide p33 was injected either i.v., s.c. or directly into the spleen of wild-type C57BL/6 mice. After 2 hours, DCs were isolated from the spleen of animals injected either i.s. or i.v., and additionally from the regional draining lymph nodes of animals injected s.c. Cells isolated from these tissues were sorted for DCs using magnetic beads coupled with a monoclonal antibody recognizing the integrin-alpha chain, a marker specific for DCs in spleen and lymph nodes. The positively and the negatively sorted cell fractions were compared regarding their capacity to *in vitro* stimulate naive CD8<sup>+</sup> T cells from TCR transgenic mice specific for LCMV-gp. Only when peptide had been directly injected into the spleen, the DC containing cell fraction stimulated CTL to proliferate, as measured by <sup>3</sup>H-thymidine uptake. This indicated that CTL induction after direct injection of peptide into lymphatic organs reflected efficient loading of DCs with peptide. In contrast, the fraction depleted for DC did not induce proliferation and DCs isolated from lymphoid organs of i.v. and s.c. injected mice were not effective stimulators. The results are shown in Figure 4.

While the present invention has been described with reference to what are presently considered to be the preferred examples, it is to be understood that the

invention is not limited to the disclosed examples. To the contrary, the invention is intended to cover various modifications and equivalent arrangements included within the spirit and scope of the appended claims.

- 5 All publications, patents and patent applications are herein incorporated by reference in their entirety to the same extent as if each individual publication, patent or patent application was specifically and individually indicated to be incorporated by reference in its entirety.

**What is claimed is:**

1. A method of inducing and/or sustaining an immunological CTL response in a mammal, which method comprises  
delivering an antigen to the mammal at a level sufficient to induce an  
5 immunologic CTL response in the mammal and  
maintaining the level of the antigen in the mammal's lymphatic system over  
time sufficient to maintain the immunologic CTL response.
2. The method of Claim 1 wherein the CTL response is maintained by delivering  
the antigen directly to the animal's lymphatic system.
- 10 3. The method of Claim 2 wherein the CTL response is maintained by delivering  
the antigen directly to the spleen, a lymph node or lymph vessel.
4. A method of treating a mammal having a disease, or being predisposed to a  
disease, to which the mammal's immune system mounts a cell-mediated response to a  
disease-related antigen to attack the disease, which method comprises  
15 delivering a disease-matched antigen to the animal at a level sufficient to  
induce an increased CTL-response in the animal and  
maintaining the increased CTL-response in the animal by sustained, regular  
delivery of the disease-matched antigen to the animal for a time sufficient to treat the  
disease wherein the sustained, regular delivery of the antigen is done in a manner that  
20 maintains the level of antigen in the animal's lymphatic system.
5. The method of Claim 4 wherein the disease is cancer.
6. The method of Claim 5 wherein the cancer is malignant melanoma.
7. The method of Claim 4 wherein the disease is an infectious disease.
8. The method of Claim 7 wherein the infectious disease is a viral disease.
- 25 9. The method of Claim 4 wherein a single antigen is delivered to the animal.
10. The method of Claim 4 wherein multiple antigens are delivered to the animal.



11. The method of Claim 4 wherein the CTL response is maintained by delivering the antigen directly to the animal's lymphatic system.
12. The method of Claim 11 wherein the CTL response is maintained by delivering the antigen directly to a lymph node or lymph vessel.
- 5 13. The method of Claim 12 wherein the antigen is delivered directly to an inguinal or axillary lymph node.
14. The method of Claim 4 wherein the antigen is delivered to the animal by pumping a physiologically-acceptable, composition of the antigen from a device held external of the animal's body through a transmission line and catheter positioned to  
10 deliver the antigen-containing composition so that the antigen reaches the animal's lymph system.
15. The method of Claim 4 wherein the antigen is delivered by implanting an implantable, sustained-release pump containing a physiologically-acceptable, composition of the antigen at or near a site of a lymphatic organ or vessel so that the  
15 antigen-containing composition is released on a sustained regular basis over time.
16. The method of Claim 15, wherein the pump is an osmotic pump.
17. The method of Claim 4, wherein the disease is cancer and the antigen is a tumor-associated antigen.
18. The method of Claim 17, wherein the antigen is selected from the group  
20 consisting of a differentiation antigen, tumor-specific multilineage antigen, an embryonic antigen, an antigen from an expressed oncogene, an antigen from an expressed mutated tumor-suppressor gene, and a viral antigen.
19. A method according to Claim 17, wherein the antigen is selected from the group consisting of MART-1/MelanA (MART-1), gp100 (Pmel 17), tyrosinase, TRP-  
25 1, TRP-2, MAGE-1, MAGE-3, BAGE, GAGE-1, GAGE-2, p15(58), CEA p53, Ras, HER-2/neu, BCR-ABL, E2A-PRL, H4-RET, IGH-IGK, MYL-RAR, EBVA, (HPV) antigens E6 and E7, TSP-180, MAGE-4, MAGE-5, MAGE-6, RAGE, NY-ESO, p185erbB2, p180erbB-3, c-met, nm-23H1, PSA, TAG-72, CA 19-9, CA 72-4, CAM 17.1, NuMa, K-ras, H-ras,  $\beta$ -Catenin, CDK4, Mum-1, p15, p16.

20. A method according to Claim 4, wherein a cytokine that is capable of enhancing the CTL response is delivered and/or maintained along with the antigen.

21. A method according to Claim 20, wherein the cytokine is GM-CSF, IL-12, IL-2, TNF, IFN $\gamma$ , IL-18, IL-3, IL-4, IL-8, IL-9, IL-13, IL-10, IL-14, IL-15, G-SCF, IFN  
5 alpha, IFN beta, IFN gamma, TGF alpha, TGF beta.

22. An article of manufacture for delivering an antigen that induces a CTL response in an animal, which article comprises

a reservoir of a physiologically-acceptable, antigen-containing composition that is capable of inducing a CTL response in an animal,

10 a pump connected to the reservoir to deliver the composition at a defined rate,

a transmission line to discharge the composition from the reservoir, and, optionally,

a delivery line connected to the transmission line, which delivery line is of a size suitable for positioning in the animal and for delivery of the composition in a  
15 manner that reaches the lymphatic system of the animal.

23. The article of Claim 22 wherein the reservoir is removable from the article of manufacture or is refillable.

24. The article of Claim 22 wherein the composition comprises only one antigen.

25. The article of Claim 22 wherein the composition comprises more than one  
20 antigen.

26. The article of Claim 22 wherein the composition further comprises a cytokine capable of enhancing a CTL response.

27. The article of Claim 26 wherein the cytokine is GM-CSF, IL-12, IL-2, TNF, IFN $\gamma$ , IL-18, IL-3, IL-4, IL-8, IL-9, IL-13, IL-10, IL-14, IL-15, G-SCF, IFN alpha,  
25 IFN beta, IFN gamma, TGF alpha, TGF beta.

28. The article of Claim 22 wherein the antigen is a differentiation antigen, a tumor-specific multilineage antigen, an embryonic antigen, an oncogene antigen, a mutated tumor-suppressor gene antigen, or a viral antigen.

29. The article of claim 28 wherein the antigen is selected from the group consisting of MART-1/MelanA (MART-1), gp100 (Pmel 17), tyrosinase, TRP-1, TRP-2, MAGE-1, MAGE-3, BAGE, GAGE-1, GAGE-2, p15(58), CEA, p53, Ras, HER-2/neu, BCR-ABL, E2A-PRL, H4-RET, IGH-IGK, MYL-RAR, EBVA, (HPV) antigens E6 and E7, TSP-180, MAGE-4, MAGE-5, MAGE-6, RAGE, NY-ESO, p185erbB2, p180erbB-3, c-met, nm-23H1, PSA, TAG-72, CA 19-9, CA 72-4, CAM 17.1, NuMa, K-ras, H-ras,  $\beta$ -Catenin, CDK4, Mum-1, p15, p16.

30. The article of Claim 22 wherein the article is an external device and the delivering line is a catheter that is long enough for delivery to the animal subcutaneously or lymphatically.

31. The article of Claim 30 wherein the catheter is long enough for delivery directly to the lymphatic system of the animal.

32. The article of Claim 31 wherein the delivery to the lymphatic system is through an axillary or inguinal node.

33. The article of Claim 22 that is portable.

34. The article of Claim 33 that is of a size suitable for portably attaching to a human.

35. The article of Claim 22 wherein the pump is a roller/peristaltic pump, a syringe pump, a piston/valve pump, or a gas pressure pump.

36. The article of Claim 22 wherein the pump is battery operated.

37. The article of Claim 22 in combination with printed instructions for delivery of the antigen composition on a regular basis over time to maintain the antigen in the animal's lymphatic system at a level sufficient to maintain a CTL response in the animal.

38. A process for preparing a system useful for inducing a sustained CTL response in an animal needing such a response, which comprises:

placing a physiologically-acceptable, aqueous, antigen-containing composition in a reservoir having a pump for delivering the composition at a defined  
5 rate through a transmission line to the animal.

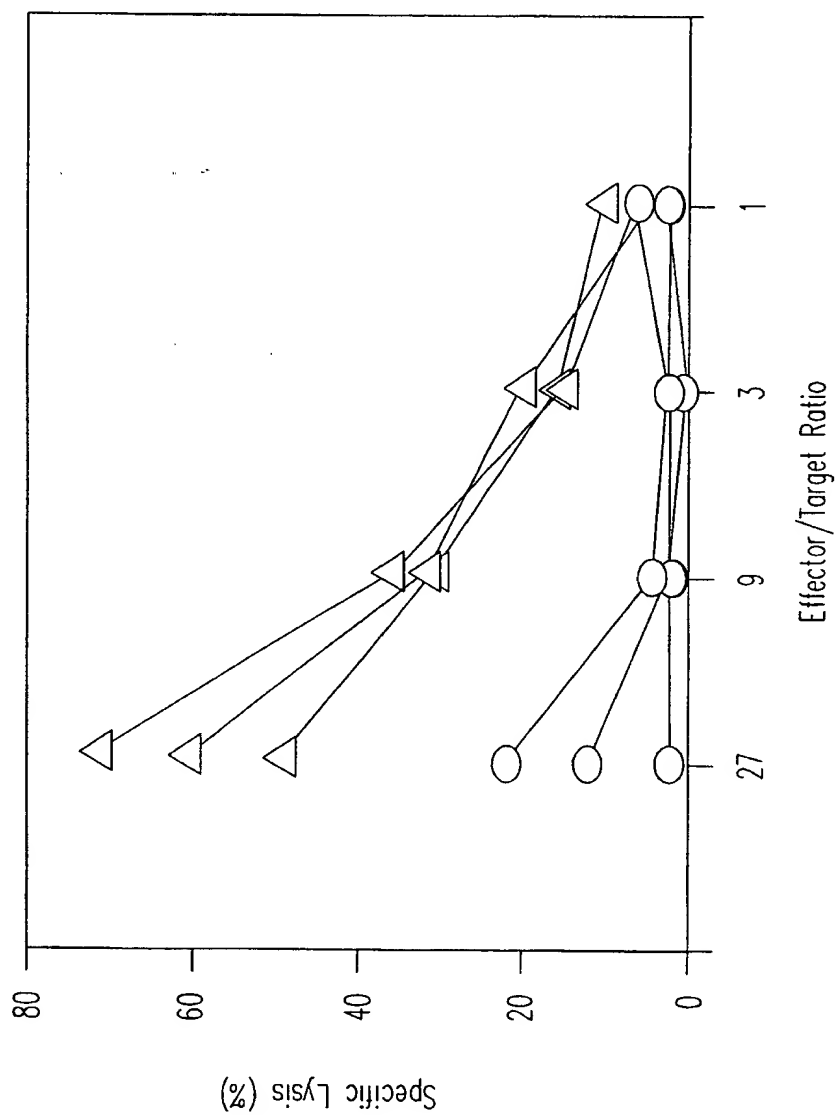


FIG. 1

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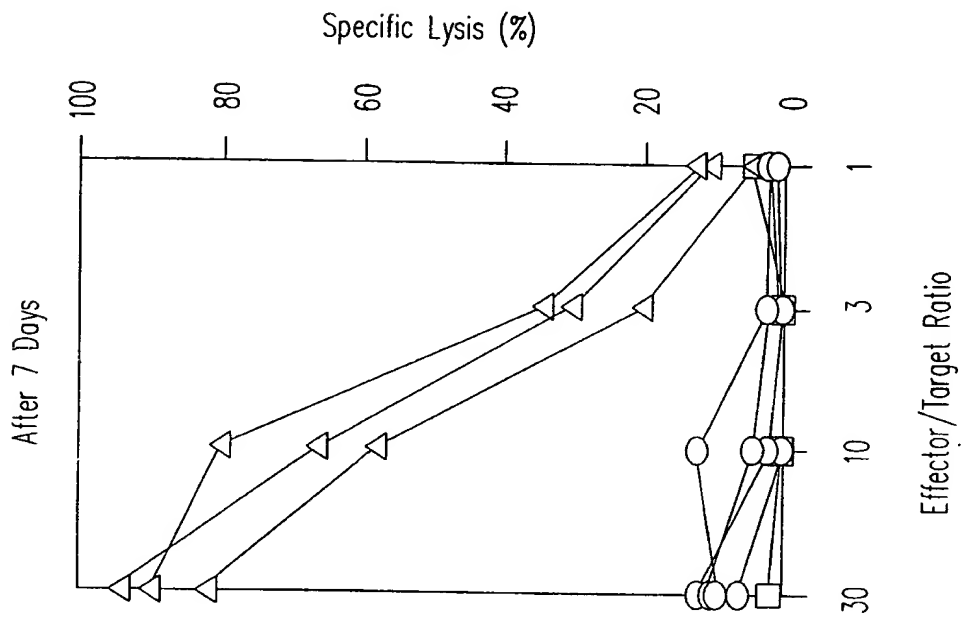


FIG. 2B

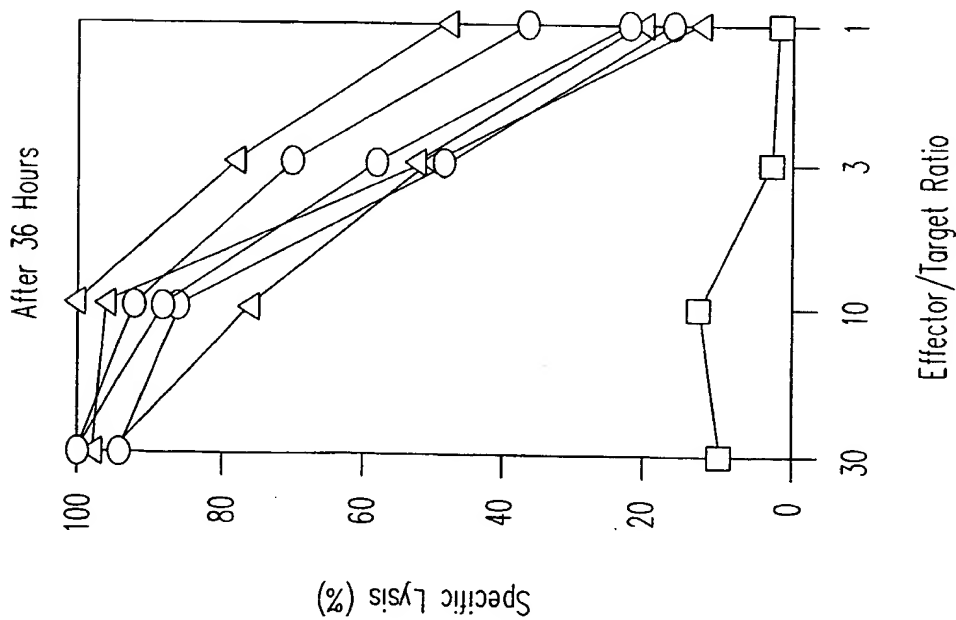


FIG. 2A

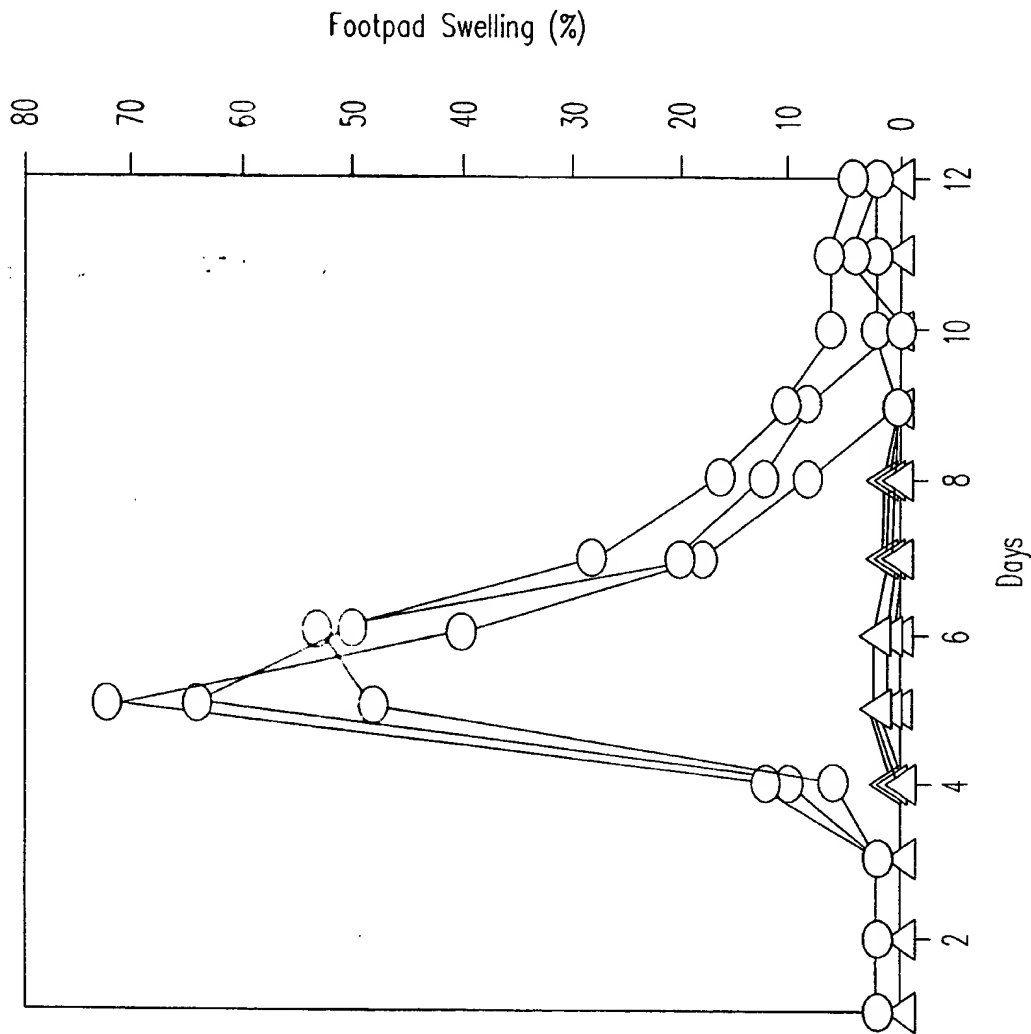


FIG. 2C

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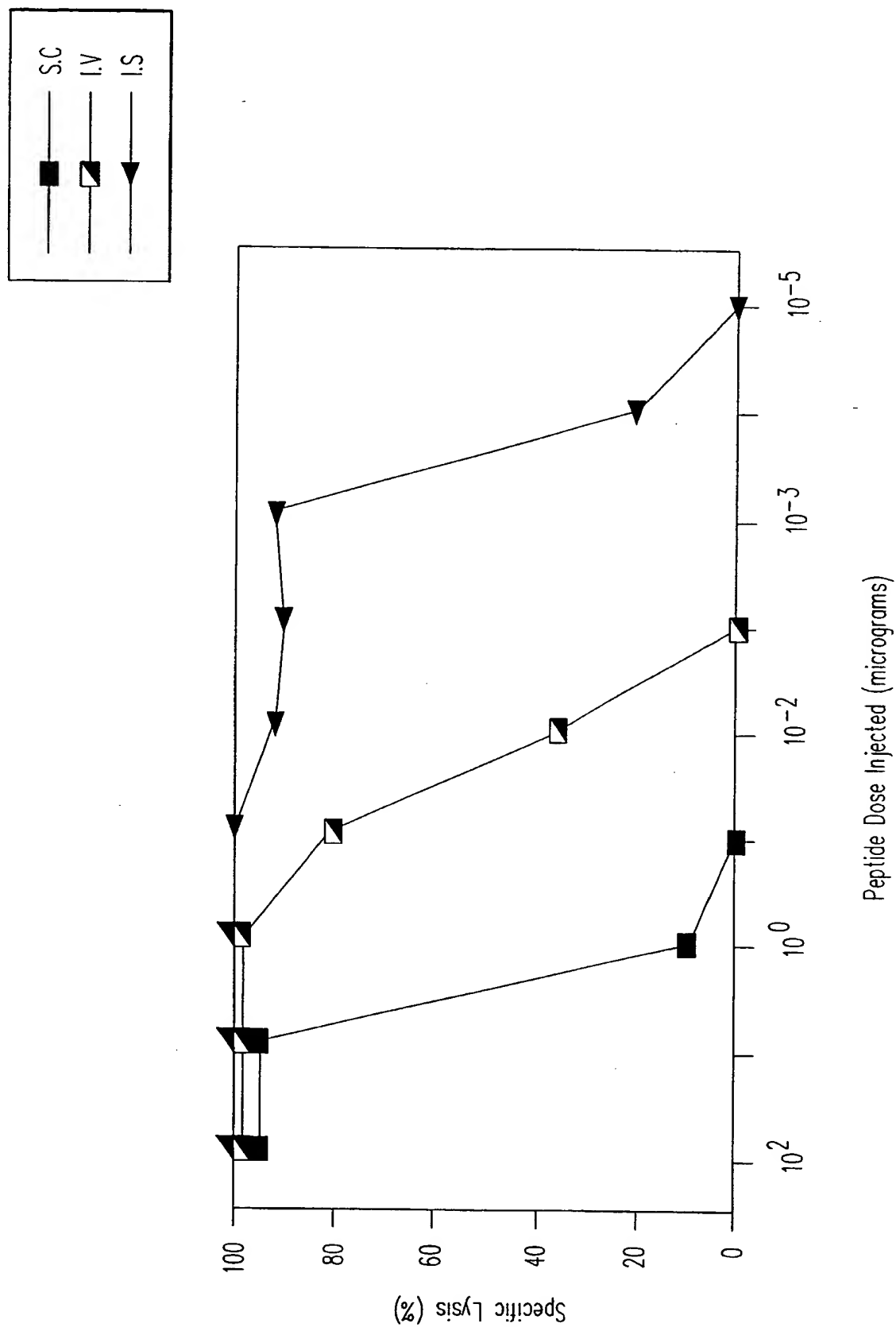
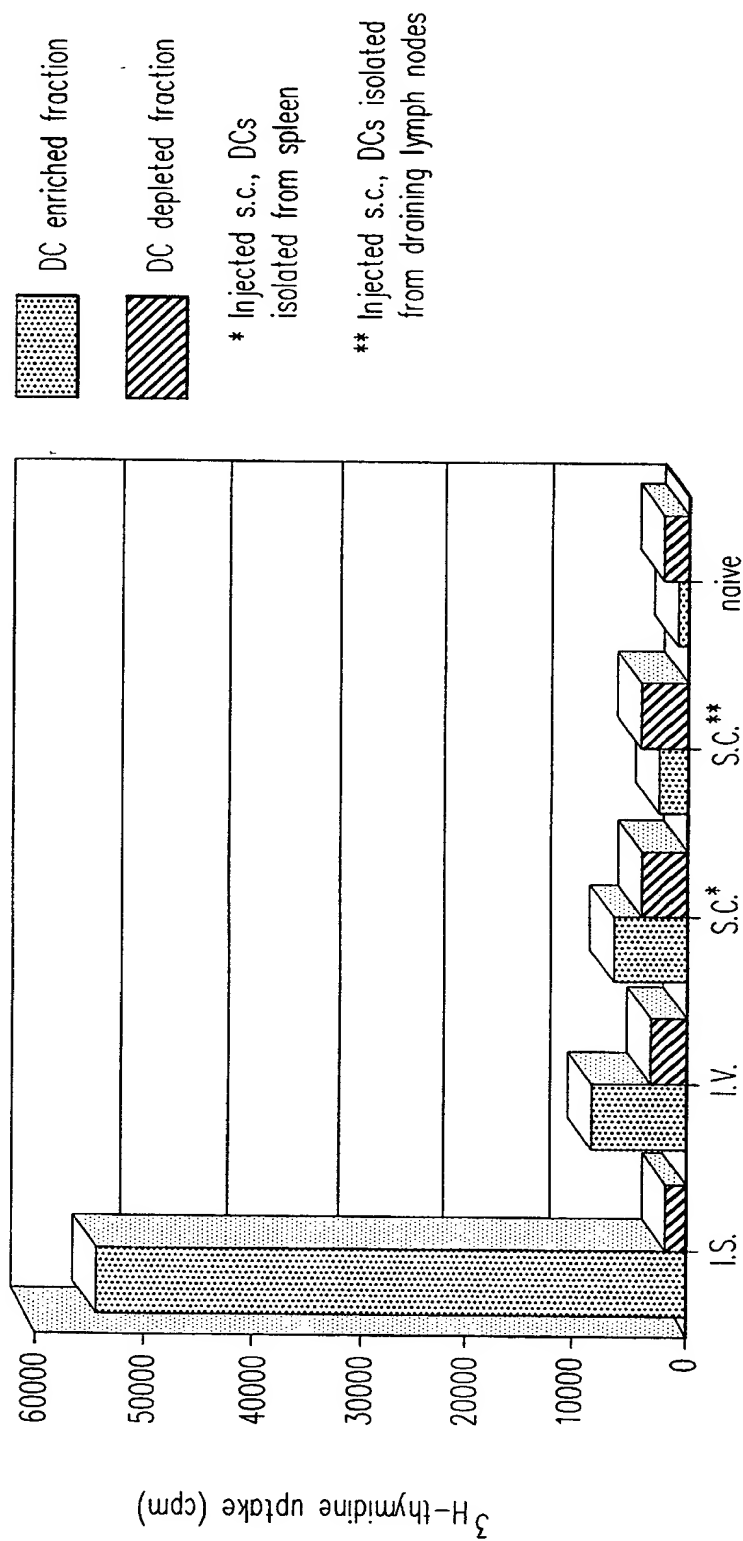


FIG. 3

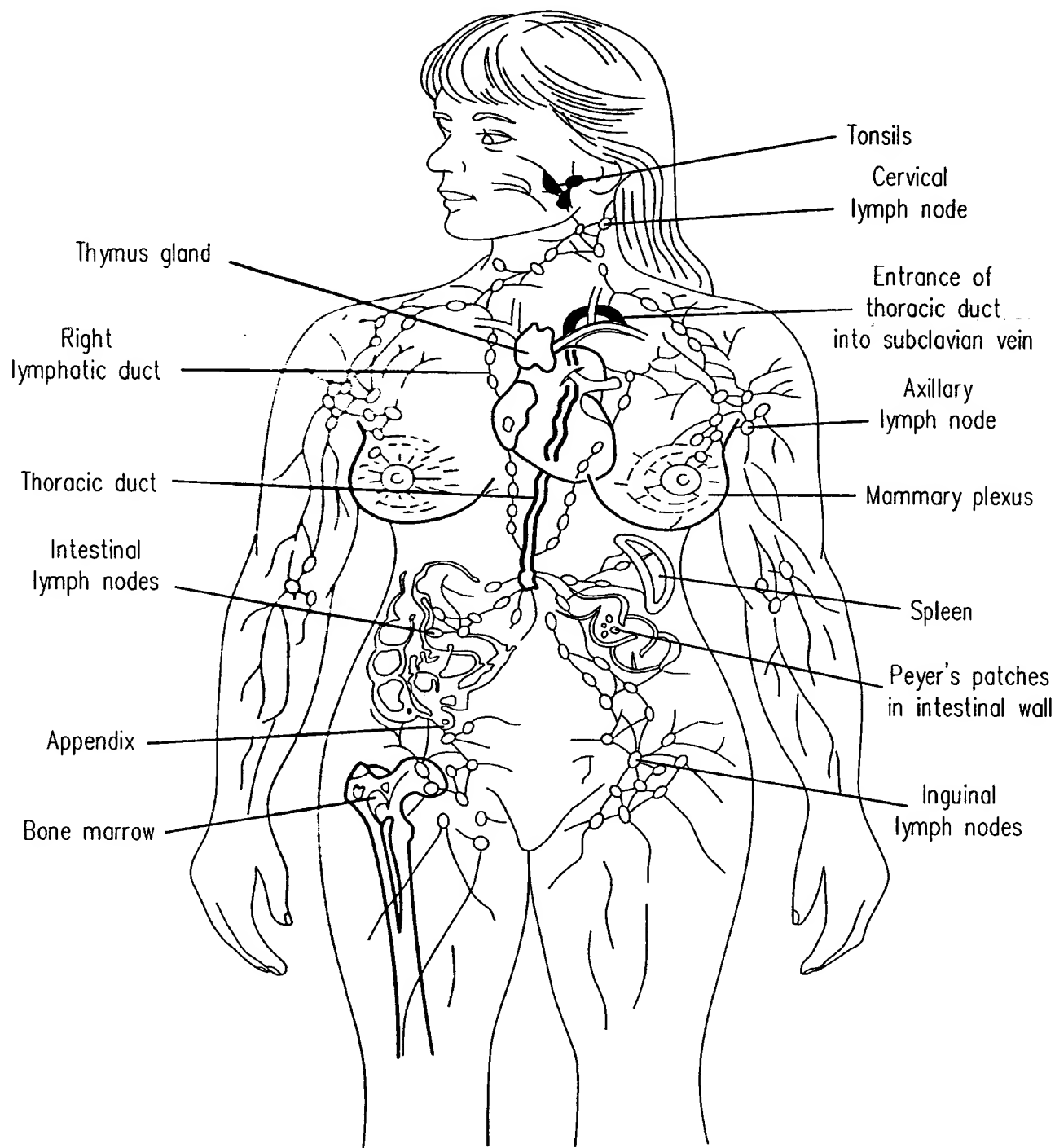




Route of Delivery

FIG. 4

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**FIG. 5**

## SEQUENCE LISTING

<110> Kuendig, Thomas M.  
Simard, John J. L.

<120> A Method of Inducing a CTL Response

<130> CTLI-001/02WO

<140> Not yet assigned

<141> 1998-07-10

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5

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10

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Ile Val Thr Asp Phe Ser Val Ile Lys

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5

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Ala Val Phe Asp Arg Lys Ser Asp Ala Lys

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Arg Tyr Ser Ile Phe Phe Asp Tyr

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5

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5

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&lt;400&gt; 39

Arg Pro Pro Ile Phe Ile Arg Arg Leu

1

5

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&lt;211&gt; 9

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Tyr Pro Leu His Glu Gln His Gly Met

1

5

&lt;210&gt; 60

&lt;211&gt; 9

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&lt;213&gt; Hepatitis C virus

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Ala Leu Met Pro Leu Tyr Ala Cys Ile

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Met Gly Leu Lys Phe Arg Gln Leu

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&lt;213&gt; Human cytomegalovirus

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&lt;213&gt; Human cytomegalovirus

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15

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&lt;211&gt; 9

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&lt;213&gt; Human cytomegalovirus

&lt;400&gt; 146

Ile Pro Ser Ile Asn Val His His Tyr

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&lt;213&gt; Human cytomegalovirus

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&lt;400&gt; 178

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Lys Ile Arg Leu Arg Pro Gly Gly Lys Lys

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Gly Arg Arg Gly Trp Glu Ala Leu Lys Tyr

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&lt;211&gt; 10

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&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 225

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&lt;210&gt; 226

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 226

Lys Ala Phe Ser Pro Glu Val Ile Pro Met Phe

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&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 227

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&lt;210&gt; 228

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

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<400> 243

Phe Pro Val Thr Pro Gln Val Pro Leu Arg  
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Thr Pro Gly Pro Gly Val Arg Tyr Pro Leu  
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Val Pro Leu Asp Glu Asp Phe Arg Lys Tyr  
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Arg Ala Ile Glu Ala Gln Ala His Leu

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&lt;210&gt; 249

&lt;211&gt; 8

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus

&lt;400&gt; 249

Thr Ala Phe Thr Ile Pro Ser Ile

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&lt;210&gt; 250

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus

&lt;400&gt; 250

Val His Pro Val His Ala Gly Pro Ile Ala

1

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&lt;210&gt; 251

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus

&lt;400&gt; 251

Asn Cys Ser Phe Asn Ile Ser Thr Ser Ile

1

5

10

&lt;210&gt; 252

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus

&lt;400&gt; 252

Cys Thr Asn Val Ser Thr Val Gln Cys

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&lt;210&gt; 253

&lt;211&gt; 9

<212> PRT

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<400> 253

Ile Gly Pro Gly Arg Ala Phe His Thr  
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<210> 254

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<400> 254

Asn Pro Asp Ile Val Ile Tyr Gln Tyr  
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<211> 10

<212> PRT

<213> Human immunodeficiency virus

<400> 255

Glu Pro Ile Val Gly Ala Glu Thr Phe Tyr  
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<211> 9

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<213> Human immunodeficiency virus

<400> 256

Glu Pro Ile Val Gly Ala Glu Thr Phe  
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<400> 257

Ser Pro Ala Ile Phe Gln Ser Ser Met  
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Val Pro Leu Asp Lys Asp Phe Arg Lys Tyr  
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Arg Pro Ile Val Ser Thr Gln Leu Leu  
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Ala Met Gln Met Leu Lys Glu Thr Ile  
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Thr Pro Tyr Asp Ile Asn Gln Met Leu  
1 5

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Arg Arg Trp Ile Gln Leu Gly Leu Gln Lys  
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&lt;400&gt; 272

Pro Leu Lys Gln His Phe Gln Ile Val

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&lt;210&gt; 273

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human papillomavirus type 11

&lt;400&gt; 273

Arg Leu Val Thr Leu Lys Asp Ile Val

1

5

&lt;210&gt; 274

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human papillomavirus type 16

&lt;400&gt; 274

Thr Leu Gly Ile Val Cys Pro Ile Cys

1

5

&lt;210&gt; 275

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human papillomavirus type 16

&lt;400&gt; 275

Gly Thr Leu Gly Ile Val Cys Pro Ile

1

5

&lt;210&gt; 276

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human papillomavirus type 16

&lt;400&gt; 276

Met Leu Asp Leu Gln Pro Glu Thr Thr

1

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&lt;210&gt; 277

&lt;211&gt; 10

<212> PRT

<213> Human papillomavirus type 16

<400> 277

Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr

1 5 10

<210> 278

<211> 8

<212> PRT

<213> Human papillomavirus type 16

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Arg Pro Arg Lys Leu Pro Gln Leu

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<213> Human papillomavirus type 16

<400> 279

Arg Ala His Tyr Asn Ile Val Thr Phe

1 5

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<212> PRT

<213> HSV

<400> 280

Ser Ser Ile Glu Phe Ala Arg Leu

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<210> 281

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<400> 281

Gly Ile Gly Ile Gly Val Leu Ala Ala

1 5

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<400> 282  
Asp Tyr Ala Thr Leu Gly Val Gly Val  
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Leu Tyr Arg Thr Phe Ala Gly Asn Pro Arg Ala  
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Gln Thr Phe Asp Phe Gly Arg Leu  
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Gly Ile Leu Gly Phe Val Phe Thr Leu  
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Lys Thr Gly Gly Pro Ile Tyr Lys Arg  
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Leu Arg Ser Arg Tyr Trp Ala Ile  
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Glu Asp Leu Arg Val Leu Ser Phe Ile  
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Gly Glu Ile Ser Pro Leu Pro Ser Leu  
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<210> 295

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Phe Glu Asp Leu Arg Val Leu Ser Phe  
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&lt;400&gt; 296

Val Ser Asp Gly Gly Pro Asn Leu Tyr

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5

&lt;210&gt; 297

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 297

Cys Thr Glu Leu Lys Leu Ser Asp Tyr

1

5

&lt;210&gt; 298

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 298

Ala Ile Met Asp Lys Asn Ile Ile Leu

1

5

&lt;210&gt; 299

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 299

Ile Met Asp Lys Asn Ile Ile Leu Lys Ala

1

5

10

&lt;210&gt; 300

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 300

Ser Arg Tyr Trp Ala Ile Arg Thr Arg

1

5

&lt;210&gt; 301

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 301

Thr Tyr Gln Arg Thr Arg Ala Leu Val

1

5

&lt;210&gt; 302

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 302

Thr Tyr Val Ser Val Ser Thr Ser Thr Leu

1

5

10

&lt;210&gt; 303

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 303

Ile Tyr Ser Thr Val Ala Ser Ser Leu

1

5

&lt;210&gt; 304

&lt;211&gt; 8

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 304

Phe Glu Ala Asn Gly Asn Leu Ile

1

5

&lt;210&gt; 305

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 305

Ile Glu Gly Gly Trp Thr Gly Met Ile

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1 5

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1 5

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1 5 10

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<400> 319  
Arg Pro Gln Ala Ser Gly Val Tyr Met  
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<210> 320  
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Phe Gln Pro Gln Asn Gly Gln Phe Ile

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&lt;210&gt; 321

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; LCMV

&lt;400&gt; 321

Ser Gly Val Glu Asn Pro Gly Gly Tyr Cys Leu

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5

10

&lt;210&gt; 322

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; LCMV

&lt;400&gt; 322

Lys Ala Val Tyr Asn Phe Ala Thr Cys Gly

1

5

10

&lt;210&gt; 323

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; MCMV

&lt;400&gt; 323

Tyr Pro His Phe Met Pro Thr Asn Leu

1

5

&lt;210&gt; 324

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; MHV

&lt;400&gt; 324

Cys Leu Ser Trp Asn Gly Pro His Leu

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&lt;210&gt; 325

&lt;211&gt; 9

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Ser Phe Ala Val Ala Thr Thr Ala Leu  
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Ser Tyr Glu Thr Phe Ile Ser Arg Leu  
1 5

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Ala Asn Tyr Asp Phe Ile Cys Val  
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Lys Ser Pro Trp Phe Thr Thr Leu  
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Ser Ser Trp Asp Phe Ile Thr Val  
1 5

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Cys Cys Leu Cys Leu Thr Val Phe Leu  
1 5

<210> 331  
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<400> 331  
Ser Pro Ser Tyr Val Tyr His Gln Phe  
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Ser Arg Arg Tyr Pro Asp Ala Val Tyr Leu His  
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<400> 333  
Arg Arg Tyr Pro Asp Ala Val Tyr Leu  
1 5

<210> 334  
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<400> 334  
Tyr Pro Ala Leu Gly Leu His Glu Phe  
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<213> Rotavirus sp.

<400> 335  
Asp Pro Val Ile Asp Arg Leu Tyr Leu  
1 5

<210> 336  
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<213> Rotavirus sp.

<400> 336  
Ser Pro Gly Arg Ser Phe Ser Tyr Phe  
1 5

<210> 337  
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<400> 337  
Thr Tyr Lys Asp Thr Val Gln Leu  
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<210> 338  
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Phe Tyr Asp Gly Phe Ser Lys Val Pro Leu  
1 5 10

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Ile Ala Gly Ile Gly Ile Leu Ala Ile  
1 5

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<400> 340  
Val Glu Ala Glu Ile Ala His Gln Ile  
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Ile Ile Tyr Arg Phe Leu Leu Ile  
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Val Gly Pro Val Phe Pro Pro Gly Met  
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Tyr Ser Gly Tyr Ile Phe Arg Asp Leu  
1 5

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&lt;400&gt; 344

Ser Tyr Ile Gly Ser Ile Asn Asn Ile

1

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&lt;210&gt; 345

&lt;211&gt; 12

&lt;212&gt; PRT

&lt;213&gt; Simian virus 40

&lt;400&gt; 345

Glu Gly Cys Thr Pro Tyr Asp Ile Asn Gln Met Leu

1

5

10

&lt;210&gt; 346

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Simian virus 40

&lt;400&gt; 346

Phe Ala Pro Gly Asn Tyr Pro Ala Leu

1

5

&lt;210&gt; 347

&lt;211&gt; 8

&lt;212&gt; PRT

&lt;213&gt; Simian virus 40

&lt;400&gt; 347

Val Val Tyr Asp Phe Leu Lys Cys

1

5

&lt;210&gt; 348

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Simian virus 40

&lt;400&gt; 348

Ser Ala Ile Asn Asn Tyr Ala Gln Lys Leu

1

5

10

&lt;210&gt; 349

&lt;211&gt; 9

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Cys Lys Gly Val Asn Lys Glu Tyr Leu

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5

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Gln Gly Ile Asn Asn Leu Asp Asn Leu

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Arg Gly Tyr Val Tyr Gln Gly Leu

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5

&lt;210&gt; 354

&lt;211&gt; 9

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&lt;400&gt; 354

Glu Ala Asp Pro Thr Gly His Ser Tyr

1

5

&lt;210&gt; 355

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 355

Val Ser Asp Gly Gly Pro Asn Leu Tyr

1

5

&lt;210&gt; 356

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 356

Cys Thr Glu Leu Lys Leu Ser Asp Tyr

1

5

&lt;210&gt; 357

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 357

Glu Val Asp Pro Ile Gly His Leu Tyr

1

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&lt;210&gt; 358

&lt;211&gt; 10

&lt;212&gt; PRT

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Met Leu Leu Ser Val Pro Leu Leu Leu Gly  
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<210> 359

<211> 9

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<213> Hepatitis B virus

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Ser Thr Asx Xaa Gln Ser Gly Xaa Gln  
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<210> 360

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<213> human

<400> 360

Tyr Met Asp Gly Thr Met Ser Gln Val  
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<210> 361

<211> 9

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<213> Human immunodeficiency virus type 1

<400> 361

Ile Leu Lys Glu Pro Val His Gly Val  
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<210> 362

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<213> Haemophilus influenzae

<400> 362

Ile Leu Gly Phe Val Phe Thr Leu Thr Val  
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Leu Leu Phe Gly Tyr Pro Val Tyr Val  
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1 5 10

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Trp Leu Ser Leu Leu Val Pro Phe Val  
1 5

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Phe Leu Pro Ser Asp Phe Phe Pro Ser Val  
1 5 10

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Cys Leu Gly Gly Leu Leu Thr Met Val  
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<213> Human cytomegalovirus

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Phe Leu Ala Gly Asn Ser Ala Tyr Glu Tyr Val  
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<400> 369

Lys Leu Gly Glu Phe Tyr Asn Gln Met Met  
1 5 10

<210> 370

<211> 10

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<213> Hepatitis C virus

<400> 370

Lys Leu Val Ala Leu Gly Ile Asn Ala Val  
1 5 10

<210> 371

<211> 9

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Asp Leu Met Gly Tyr Ile Pro Leu Val  
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<210> 372

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<213> Human papillomavirus type 11

<400> 372

Arg Leu Val Thr Leu Lys Asp Ile Val



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&lt;210&gt; 373

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 373

Met Leu Leu Ala Val Leu Tyr Cys Leu

1

5

&lt;210&gt; 374

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 374

Ala Ala Gly Ile Gly Ile Leu Thr Val

1

5

&lt;210&gt; 375

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 375

Tyr Leu Glu Pro Gly Pro Val Thr Ala

1

5

&lt;210&gt; 376

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 376

Ile Leu Asp Gly Thr Ala Thr Leu Arg Leu

1

5

10

&lt;210&gt; 377

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 377

Leu Leu Asp Gly Thr Ala Thr Leu Arg Leu  
1 5 10

&lt;210&gt; 378

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 378

Ile Thr Asp Gln Val Pro Phe Ser Val  
1 5

&lt;210&gt; 379

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 379

Lys Thr Trp Gly Gln Tyr Trp Gln Val  
1 5

&lt;210&gt; 380

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 380

Thr Ile Thr Asp Gln Val Pro Phe Ser Val  
1 5 10

&lt;210&gt; 381

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 381

Ala Phe His His Val Ala Arg Glu Leu  
1 5

&lt;210&gt; 382

&lt;211&gt; 9

&lt;212&gt; PRT

<213> Plasmodium falciparum

<400> 382

Tyr Leu Asn Lys Ile Gln Asn Ser Leu  
1 5

<210> 383

<211> 10

<212> PRT

<213> Plasmodium falciparum

<400> 383

Met Met Arg Lys Leu Ala Ile Leu Ser Val  
1 5 10

<210> 384

<211> 10

<212> PRT

<213> Haemophilus influenzae

<400> 384

Lys Ala Gly Glu Phe Tyr Asn Gln Met Met  
1 5 10

<210> 385

<211> 9

<212> PRT

<213> EBV

<400> 385

Asn Ile Ala Glu Gly Leu Arg Ala Leu  
1 5

<210> 386

<211> 9

<212> PRT

<213> EBV

<400> 386

Asn Leu Arg Arg Gly Thr Ala Leu Ala  
1 5

<210> 387

<211> 9  
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<400> 387  
Ala Leu Ala Ile Pro Gln Cys Arg Leu  
1 5

<210> 388  
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<400> 388  
Val Leu Lys Asp Ala Ile Lys Asp Leu  
1 5

<210> 389  
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<400> 389  
Phe Met Val Phe Ile Gln Thr His Ile  
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<400> 390  
His Leu Ile Val Asp Thr Asp Ser Leu  
1 5

<210> 391  
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<400> 391  
Ser Leu Gly Asn Pro Ser Leu Ser Val  
1 5

<210> 392

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<213> EBV

<400> 392

Pro Leu Ala Ser Ala Met Arg Met Leu

1

5

<210> 393

<211> 9

<212> PRT

<213> EBV

<400> 393

Arg Met Leu Trp Met Ala Asn Tyr Ile

1

5

<210> 394

<211> 9

<212> PRT

<213> EBV

<400> 394

Met Leu Trp Met Ala Asn Tyr Ile Val

1

5

<210> 395

<211> 9

<212> PRT

<213> EBV

<400> 395

Ile Leu Pro Gln Gly Pro Gln Thr Ala

1

5

<210> 396

<211> 9

<212> PRT

<213> EBV

<400> 396

Pro Leu Arg Pro Thr Ala Pro Thr Ile

1

5

&lt;210&gt; 397

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 397

Pro Leu Pro Pro Ala Thr Leu Thr Val

1

5

&lt;210&gt; 398

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 398

Arg Met His Leu Pro Val Leu His Val

1

5

&lt;210&gt; 399

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 399

Pro Met Pro Leu Pro Pro Ser Gln Leu

1

5

&lt;210&gt; 400

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 400

Gln Leu Pro Pro Pro Ala Ala Pro Ala

1

5

&lt;210&gt; 401

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 401

Ser Met Pro Glu Leu Ser Pro Val Leu

1

5

&lt;210&gt; 402

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 402

Asp Leu Asp Glu Ser Trp Asp Tyr Ile

1

5

&lt;210&gt; 403

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 403

Pro Leu Pro Cys Val Leu Trp Pro Val

1

5

&lt;210&gt; 404

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 404

Ser Leu Glu Glu Cys Asp Ser Glu Leu

1

5

&lt;210&gt; 405

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 405

Glu Ile Lys Arg Tyr Lys Asn Arg Val

1

5

&lt;210&gt; 406

&lt;211&gt; 9

&lt;212&gt; PRT

<213> EBV

<400> 406

Gln Leu Leu Gln His Tyr Arg Glu Val  
1 5

<210> 407

<211> 9

<212> PRT

<213> EBV

<400> 407

Leu Leu Gln His Tyr Arg Glu Val Ala  
1 5

<210> 408

<211> 9

<212> PRT

<213> EBV

<400> 408

Leu Leu Lys Gln Met Cys Pro Ser Leu  
1 5

<210> 409

<211> 9

<212> PRT

<213> EBV

<400> 409

Ser Ile Ile Pro Arg Thr Pro Asp Val  
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<210> 410

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<213> Haemophilus influenzae

<400> 410

Ala Ile Met Asp Lys Asn Ile Ile Leu  
1 5

<210> 411



<211> 10  
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<400> 411  
Ile Met Asp Lys Asn Ile Ile Leu Lys Ala  
1 5 10

<210> 412  
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<400> 412  
Leu Leu Ala Leu Leu Ser Cys Leu Thr Val  
1 5 10

<210> 413  
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<212> PRT  
<213> Hepatitis C virus

<400> 413  
Ile Leu His Thr Pro Gly Cys Val  
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<210> 414  
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Gln Leu Arg Arg His Ile Asp Leu Leu Val  
1 5 10

<210> 415  
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<400> 415  
Asp Leu Cys Gly Ser Val Phe Leu Val  
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Ser Met Val Gly Asn Trp Ala Lys Val  
1 5

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His Leu His Gln Asn Ile Val Asp Val  
1 5

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Phe Leu Leu Leu Ala Asp Ala Arg Val  
1 5

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<400> 419  
Gly Leu Arg Asp Leu Ala Val Ala Val Glu Pro Val Val  
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<210> 420  
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<212> PRT  
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Ser Leu Leu Ala Pro Gly Ala Lys Gln Asn Val

1 5 10

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<400> 421  
Leu Leu Ala Pro Gly Ala Lys Gln Asn Val  
1 5 10

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Phe Leu Leu Ser Leu Gly Ile His Leu  
1 5

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<400> 423  
Ser Leu Tyr Ala Asp Ser Pro Ser Val  
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<400> 424  
Gly Leu Ser Arg Tyr Val Ala Arg Leu  
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<210> 425  
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<213> human

&lt;400&gt; 425

Lys Ile Phe Gly Ser Leu Ala Phe Leu  
1 5

&lt;210&gt; 426

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 426

Glu Leu Val Ser Glu Phe Ser Arg Met  
1 5

&lt;210&gt; 427

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 427

Lys Leu Thr Pro Leu Cys Val Thr Leu  
1 5

&lt;210&gt; 428

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 428

Ser Leu Leu Asn Ala Thr Asp Ile Ala Val  
1 5 10

&lt;210&gt; 429

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 429

Val Leu Tyr Arg Tyr Gly Ser Phe Ser Val  
1 5 10

&lt;210&gt; 430

&lt;211&gt; 9

&lt;212&gt; PRT

<213> human

<400> 430

Tyr Ile Gly Glu Val Leu Val Ser Val  
1 5

<210> 431

<211> 10

<212> PRT

<213> Hepatitis C virus

<400> 431

Leu Leu Phe Asn Ile Leu Gly Gly Trp Val  
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<210> 432

<211> 10

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<213> Hepatitis B virus

<400> 432

Leu Leu Val Pro Phe Val Gln Trp Phe Trp  
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<210> 433

<211> 9

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<213> Hepatitis B virus

<400> 433

Ala Leu Met Pro Leu Tyr Ala Cys Ile  
1 5

<210> 434

<211> 9

<212> PRT

<213> Hepatitis C virus

<400> 434

Tyr Leu Val Ala Tyr Gln Ala Thr Val  
1 5

<210> 435

<211> 9

<212> PRT

<213> Human papillomavirus type 16

<400> 435

Thr Leu Gly Ile Val Cys Pro Ile Cys

1

5

<210> 436

<211> 10

<212> PRT

<213> Hepatitis C virus

<400> 436

Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu

1

5

10

<210> 437

<211> 10

<212> PRT

<213> Hepatitis B virus

<400> 437

Leu Leu Pro Ile Phe Phe Cys Leu Trp Val

1

5

10

<210> 438

<211> 9

<212> PRT

<213> Hepatitis B virus

<400> 438

Tyr Met Asp Asp Val Val Leu Gly Ala

1

5

<210> 439

<211> 9

<212> PRT

<213> Human papillomavirus type 16

<400> 439

Gly Thr Leu Gly Ile Val Cys Pro Ile

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<400> 440  
Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile  
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<210> 441  
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Met Leu Asp Leu Gln Pro Glu Thr Thr  
1 5

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<400> 442  
Ser Leu Met Ala Phe Thr Ala Ala Val  
1 5

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Cys Ile Asn Gly Val Cys Trp Thr Val  
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Val Met Asn Ile Leu Leu Gln Tyr Val Val

1 5 10

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Ile Leu Thr Val Ile Leu Gly Val Leu  
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Phe Leu Trp Gly Pro Arg Ala Leu Val  
1 5

<210> 447  
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<400> 447  
Leu Leu Cys Pro Ala Gly His Ala Val  
1 5

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<400> 448  
Ile Leu Asp Ser Phe Asp Pro Leu Val  
1 5

<210> 449  
<211> 9  
<212> PRT  
<213> Hepatitis B virus



&lt;400&gt; 449

Leu Leu Leu Cys Leu Ile Phe Leu Leu

1

5

&lt;210&gt; 450

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Hepatitis B virus

&lt;400&gt; 450

Leu Ile Asp Tyr Gln Gly Met Leu Pro Val

1

5

10

&lt;210&gt; 451

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Hepatitis B virus

&lt;400&gt; 451

Ser Ile Val Ser Pro Phe Ile Pro Leu Leu

1

5

10

&lt;210&gt; 452

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Hepatitis B virus

&lt;400&gt; 452

Phe Leu Leu Thr Arg Ile Leu Thr Ile

1

5

&lt;210&gt; 453

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Plasmodium falciparum

&lt;400&gt; 453

His Leu Gly Asn Val Lys Tyr Leu Val

1

5

&lt;210&gt; 454

&lt;211&gt; 9

&lt;212&gt; PRT

<213> Plasmodium falciparum

<400> 454

Gly Ile Ala Gly Gly Leu Ala Leu Leu  
1 5

<210> 455

<211> 9

<212> PRT

<213> Hepatitis C virus

<400> 455

Ile Leu Ala Gly Tyr Gly Ala Gly Val  
1 5

<210> 456

<211> 9

<212> PRT

<213> Hepatitis C virus

<400> 456

Gly Leu Gln Asp Cys Thr Met Leu Val  
1 5

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<211> 10

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<213> Hepatitis C virus

<400> 457

Thr Gly Ala Pro Val Thr Tyr Ser Thr Tyr  
1 5 10

<210> 458

<211> 9

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<213> Human immunodeficiency virus type 1

<400> 458

Val Ile Tyr Gln Tyr Met Asp Asp Leu  
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<210> 459

<211> 10  
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<400> 459  
Val Leu Pro Asp Val Phe Ile Arg Cys Val  
1 5 10

<210> 460  
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Val Leu Pro Asp Val Phe Ile Arg Cys  
1 5

<210> 461  
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<400> 461  
Ala Val Gly Ile Gly Ile Ala Val Val  
1 5

<210> 462  
<211> 9  
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<400> 462  
Leu Val Val Leu Gly Leu Leu Ala Val  
1 5

<210> 463  
<211> 9  
<212> PRT  
<213> human

<400> 463  
Ala Leu Gly Leu Gly Leu Leu Pro Val  
1 5

<210> 464  
<211> 9  
<212> PRT  
<213> HSV-2

<400> 464  
Gly Ala Gly Ile Gly Val Ala Val Leu  
1 5

<210> 465  
<211> 9  
<212> PRT  
<213> Pseudorabies virus

<400> 465  
Ile Ala Gly Ile Gly Ile Leu Ala Ile  
1 5

<210> 466  
<211> 9  
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<213> Adenovirus 3

<400> 466  
Leu Ile Val Ile Gly Ile Leu Ile Leu  
1 5

<210> 467  
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<213> S. lincolnensis

<400> 467  
Leu Ala Gly Ile Gly Leu Ile Ala Ala  
1 5

<210> 468  
<211> 9  
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<213> yeast

<400> 468  
Val Asp Gly Ile Gly Ile Leu Thr Ile

1 5

<210> 469

<211> 9

<212> PRT

<213> B. polymyxa

<400> 469

Gly Ala Gly Ile Gly Val Leu Thr Ala

1 5

<210> 470

<211> 9

<212> PRT

<213> Escherichia coli

<400> 470

Ala Ala Gly Ile Gly Ile Ile Gln Ile

1 5

<210> 471

<211> 9

<212> PRT

<213> Escherichia coli

<400> 471

Gln Ala Gly Ile Gly Ile Leu Leu Ala

1 5

<210> 472

<211> 11

<212> PRT

<213> human

<400> 472

Lys Ala Arg Asp Pro His Ser Gly His Phe Val

1 5 10

<210> 473

<211> 11

<212> PRT

<213> human

&lt;400&gt; 473

Lys Ala Cys Asp Pro His Ser Gly His Phe Val

1 5 10

&lt;210&gt; 474

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 474

Ala Cys Asp Pro His Ser Gly His Phe Val

1 5 10

&lt;210&gt; 475

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 475

Ser Leu Tyr Asn Thr Val Ala Thr Leu

1 5

&lt;210&gt; 476

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 476

Glu Leu Val Ser Glu Phe Ser Arg Val

1 5

&lt;210&gt; 477

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 477

Arg Gly Pro Gly Arg Ala Phe Val Thr Ile

1 5 10

&lt;210&gt; 478

&lt;211&gt; 9

&lt;212&gt; PRT

<213> Hepatitis C virus

<400> 478

His Met Trp Asn Phe Ile Ser Gly Ile

1

5

<210> 479

<211> 10

<212> PRT

<213> Human cytomegalovirus

<400> 479

Asn Leu Val Pro Met Val Ala Thr Val Gln

1

5

10

<210> 480

<211> 9

<212> PRT

<213> Human papillomavirus type 6b

<400> 480

<210> 481

<211> 9

<212> PRT

<213> Human papillomavirus type 6b

<400> 481

Pro Leu Lys Gln His Phe Gln Ile Val

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<210> 482

<211> 10

<212> PRT

<213> EBV

<400> 482

Leu Leu Asp Phe Val Arg Phe Met Gly Val

1

5

10

<210> 483

<211> 9

<212> PRT

<213> Haemophilus influenzae

<400> 483

Ala Ile Met Glu Lys Asn Ile Met Leu

1

5

<210> 484

<211> 9

<212> PRT

<213> Plasmodium falciparum

<400> 484

Tyr Leu Lys Thr Ile Gln Asn Ser Leu

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<211> 9

<212> PRT

<213> Plasmodium falciparum

<400> 485

Tyr Leu Asn Lys Ile Gln Asn Ser Leu

1

5

<210> 486

<211> 10

<212> PRT

<213> Human papillomavirus type 16

<400> 486

Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr

1

5

10

<210> 487

<211> 9

<212> PRT

<213> Human papillomavirus type 16

<400> 487

Leu Leu Met Gly Thr Leu Gly Ile Val

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<400> 488  
Thr Leu Gly Ile Val Cys Pro Ile  
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Lys Leu Pro Gln Leu Cys Thr Glu Leu  
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<400> 491  
Thr Ile His Asp Ile Ile Leu Glu Cys  
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Leu Gly Ile Val Cys Pro Ile Cys Ser  
1 5

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<400> 493  
Val Ile Leu Gly Val Leu Leu Leu Ile  
1 5

<210> 494  
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<213> 'Axial Seamount' polynoid polychaete

<400> 494  
Ala Leu Met Asp Lys Ser Leu His Val  
1 5

<210> 495  
<211> 9  
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<400> 495  
Gly Ile Leu Thr Val Ile Leu Gly Val  
1 5

<210> 496  
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<400> 496  
Met Ile Asn Ala Tyr Leu Asp Lys Leu  
1 5

<210> 497  
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<400> 497  
Ala Ala Gly Ile Gly Ile Leu Thr Val

1

5

&lt;210&gt; 498

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 498

Phe Leu Pro Ser Asp Phe Phe Pro Ser Val

1

5

10

&lt;210&gt; 499

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; EBV

&lt;400&gt; 499

Ser Val Arg Asp Arg Leu Ala Arg Leu

1

5

&lt;210&gt; 500

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 500

Phe Ala Tyr Asp Gly Lys Asp Tyr Ile

1

5

&lt;210&gt; 501

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 501

Ala Leu Leu Ala Val Gly Ala Thr Lys

1

5

&lt;210&gt; 502

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 502

Arg Leu Arg Asp Leu Leu Leu Ile Val Thr Arg  
1 5 10

&lt;210&gt; 503

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 503

Gln Val Pro Leu Arg Pro Met Thr Tyr Lys  
1 5 10

&lt;210&gt; 504

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 504

Thr Val Tyr Tyr Gly Val Pro Val Trp Lys  
1 5 10

&lt;210&gt; 505

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Human immunodeficiency virus type 1

&lt;400&gt; 505

Arg Leu Arg Pro Gly Gly Lys Lys Lys  
1 5

&lt;210&gt; 506

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Haemophilus influenzae

&lt;400&gt; 506

Ile Leu Arg Gly Ser Val Ala His Lys  
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&lt;210&gt; 507

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&lt;400&gt; 552

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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

|   |  |           |  |
|---|--|-----------|--|
| <b>(51) International Patent Classification <sup>6</sup> :</b><br><b>A61K 39/00, 38/19, 9/00 // (A61K 39/00, 38:19)</b>   |  | <b>A3</b> | <b>(11) International Publication Number:</b> <b>WO 99/02183</b><br><b>(43) International Publication Date:</b> 21 January 1999 (21.01.99)   |
| <b>(21) International Application Number:</b> PCT/US98/14289<br><b>(22) International Filing Date:</b> 10 July 1998 (10.07.98)<br><b>(30) Priority Data:</b><br>2,209,815 10 July 1997 (10.07.97) CA<br>08/988,320 10 December 1997 (10.12.97) US<br><b>(63) Related by Continuation (CON) or Continuation-in-Part (CIP) to Earlier Application</b><br>US 08/988,320 (CIP)<br>Filed on 10 December 1997 (10.12.97)<br><b>(71) Applicant (for all designated States except US):</b> CTL IMMUNOTHERAPIES CORPORATION [CA/CA]; Suite 5100, One First Canadian Place, Toronto, Ontario M5X 1K2 (CA).<br><b>(72) Inventors; and</b><br><b>(75) Inventors/Applicants (for US only):</b> KÜNDIG, Thomas, M. [US/CH]; Schwamendingen Strasse #79, CH-8050 Zurich (CH). SIMARD, John, J., L. [-/CA]; Suite 1014, 24 Wellesley Street, West, Toronto, Ontario M4Y 1G1 (CA). |  |           | <b>(74) Agents:</b> MORAN, Tom, M.; Cooley Godward LLP, 3000 El Camino Real, Five Palo Alto Square, Palo Alto, CA 94306-2155 (US) et al.<br><b>(81) Designated States:</b> AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, GM, HR, HU, ID, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).<br><b>Published</b><br><i>With international search report.</i><br><i>Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i><br><b>(88) Date of publication of the international search report:</b> 14 May 1999 (14.05.99) |
| <b>(54) Title:</b> A METHOD OF INDUCING A CTL RESPONSE<br><b>(57) Abstract</b><br><p>A method of inducing a cytotoxic T-lymphocyte (CTL) response to an antigen is disclosed. The method involves delivering the antigen to the lymphatic system of an animal regularly over a sustained period of time using, e.g., an osmotic pump. The method is advantageous over prior art methods for inducing a CTL response in that it does not require repetitive immunizations or the use of adjuvants. The method of the present invention can be used for the induction of CTLs in tumor or infectious disease immunotherapy.</p>   |  |           |  |

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| EE | Estonia                  | LR | Liberia                                  | SG | Singapore                                    |    |                          |

# INTERNATIONAL SEARCH REPORT

International Application No

PCT/US 98/14289

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 6 A61K39/00 A61K38/19 A61K9/00 //(A61K39/00, 38:19)

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

| Category | Citation of document, with indication, where appropriate, of the relevant passages  | Relevant to claim No.                    |
|----------|---|--|
| X        | W0 95 17167 A (GFF) 29 June 1995<br>see the whole document  | 1,4-10,<br>17-19                         |
| Y        | R. PETERS ET AL.: "Tryptophan and serotonin metabolism after sustained tryptophan infusion."<br>NEUROCHEMISTRY INTERNATIONAL,<br>vol. 6, no. 5, 1984, pages 685-692,<br>XP002095831<br>see abstract | 1,4,7-9,<br>14-16,<br>22-24,<br>28,30-38 |

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

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Date of the actual completion of the international search

8 March 1999

Date of mailing of the international search report

29/03/1999

Name and mailing address of the ISA

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Fax: (+31-70) 340-3016

Authorized officer

Nooij, F

## INTERNATIONAL SEARCH REPORT

International Application No

PCT/US 98/14289

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

| Category | Citation of document, with indication, where appropriate, of the relevant passages   | Relevant to claim No.                                 |
|----------|--|---|
| Y        | <p>S. OEHEN ET AL.: "Antivirally protective cytotoxic T cell memory to lymphocytic choriomeningitis virus is governed by persisting antigen."<br/> THE JOURNAL OF EXPERIMENTAL MEDICINE,<br/> vol. 176, no. 5, 1 November 1992, pages 1273-1281, XP002095832<br/> New York, NY, USA<br/> see abstract<br/> see page 1278, left-hand column, line 33 - line 36<br/> see page 1279, right-hand column, line 18 - line 33</p> | <p>1,4,7-9,<br/> 14-16,<br/> 22-24,<br/> 28,30-38</p> |
| A        | <p>M. OLDSTONE ET AL.: "Discriminated selection among viral peptides with the appropriate anchor residues: Implications for the size of the cytotoxic T-lymphocyte repertoire and control of viral infection."<br/> JOURNAL OF VIROLOGY,<br/> vol. 69, no. 12, December 1995, pages 7423-7429, XP002095833<br/> Washington, DC, USA<br/> see abstract</p>  | <p>1-38</p>   |
| A        | <p>T. KÜNDIG ET AL.: "On the role of antigen in maintaining cytotoxic T-cell memory."<br/> PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE U.S.A.,<br/> vol. 93, no. 18, 3 September 1996, pages 9716-9723, XP002095834<br/> Washington, DC, USA<br/> see abstract</p>  | <p>1-38</p>   |
| A        | <p>D. MOSKOPHIDIS ET AL.: "Immunobiology of cytotoxic T-cell escape mutants of lymphocytic choriomeningitis virus."<br/> JOURNAL OF VIROLOGY,<br/> vol. 69, no. 4, April 1995, pages 2187-2193, XP002095835<br/> Washington, DC, USA<br/> see abstract<br/> see figure 1A</p>  | <p>1-38</p>   |

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# INTERNATIONAL SEARCH REPORT

International Application No

PCT/US 98/14289

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

| Category | Citation of document, with indication, where appropriate, of the relevant passages  | Relevant to claim No.         |
|----------|---|-------------------------------|
| P,X      | <p>L. DURRANT: "Cancer vaccines."<br/> ANTI-CANCER DRUGS,<br/> vol. 8, no. 8, September 1997, pages<br/> 727-733, XP002095836<br/> Oxford, GB<br/> see page 728, right-hand column, line 1 -<br/> line 36<br/> see page 730, left-hand column, line 42 -<br/> right-hand column, line 8<br/> see page 730, right-hand column, line 19 -<br/> line 40</p> <p>-----</p> | <p>1,4-6,9,<br/> 10,17-21</p> |

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/US 98/14289

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:  
Remark: Although claims 1-21 are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.
2. ☐ Claims Nos.:  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

### Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

# INTERNATIONAL SEARCH REPORT

information on patent family members

International Application No

PCT/US 98/14289

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